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**A clinical and serological study of adult and juvenile idiopathic inflammatory myopathy**

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# **A CLINICAL AND SEROLOGICAL STUDY OF ADULT AND JUVENILE IDIOPATHIC INFLAMMATORY MYOPATHY**

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**A thesis submitted for the degree of Doctor of Philosophy**

**University of Bath  
School for Health**

**September 2010**

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*For my mother*



## **GLOSSARY**

### **List of abbreviations**

#### **A. Methods**

AMP	Ammonium persulphate
BS3	Bis(sulphosuccinimidyl)suberate
dFCS	dialysed Foetal Calf Serum
DMP	Dimethyl pimelimidate-2 HCL
DMSO	Dimethyl sulfoxide
EDTA	Ethylenediaminetetraacetic acid
ELISA	Enzyme-linked immunoabsorbant assay
FCS	Foetal Calf Serum
FITC	Fluorescein isothiocyanate
HCL	Hydrogen chloride
HEp-2	Human epithelial cell line
IB	Immunoblotting
IIF	Indirect immunofluorescence
IPP	Immunoprecipitation
K562	Human chronic myelogenous leukaemic cell line
MALDI-TOF	Matrix-assisted laser desorption/ionisation-time of flight mass spectrometry
MBq	Megabecquerel
NaCl	Sodium Chloride
NaHCO <sub>3</sub>	Sodium Bicarbonate
PBS	Phosphate-buffered Saline
RPMI	Rosewell Park Memorial Institute 1640 medium
SDS-PAGE	Sodium dodecyl sulphate polyacrylamide gel electrophoresis
TBS	Tris-buffered Saline
TEMED	N,N,N <sup>1</sup> ,N <sup>1</sup> -tetramethylethylenediamine
Tris	Tris[hydroxymethyl]aminomethane
WB	Western Blotting

## **GLOSSARY**

### **List of abbreviations**

#### **B. Clinical**

AIP	Acute interstitial pneumonia
ANA	Anti-nuclear autoantibody
ANoA	Anti-nucleolar autoantibody
AOMIC	Adult Onset Immunogenetic Collaboration
ARS	Aminoacyl-tRNA
ASS	Anti-synthetase syndrome synthetases
ATP	Adenosine-5'-triphosphate
ATS	American Thoracic Society
BIRD	Bath Institute for Rheumatic Diseases
CADM	Clinically-amyopathic dermatomyositis
CAT	Cutaneous Assessment Tool
CHAQ	Childhood Health Assessment Questionnaire
CK	Creatinine kinase
CMAS	Childhood Myositis Assessment Scale
COP	Organising pneumonia
CRP	C-reactive protein
CTD	Connective Tissue Disease
DM	Dermatomyositis
DNA	Deoxyribonucleic acid
DPLD	Diffuse parenchymal lung disease
EMG	Electromyogram
ENA	Extractable nuclear antigen
ER	Endoplasmic reticulum
ERS	European Respiratory Society
ESR	Erythrocyte sedimentation rate
ET-1	Endothelin-1
GI	Gastrointestinal
HLA	Human leucocyte antigen
HRCT	High-resolution computerised tomography
IBM	Inclusion body myositis
ICAM-1	Intracellular adhesion molecule-1
IFN	Interferon
IL	Interleukin
IIM	Idiopathic inflammatory myopathy

## **GLOSSARY**

### **List of abbreviations**

#### **B. Clinical**

JDM	Juvenile dermatomyositis
JDRR	Juvenile dermatomyositis Registry and Repository
LDH	Lactate dehydrogenase
MAA	Myositis-associated autoantibody
MCTD	Mixed connective tissue disease
MDA5	Melanoma-differentiation associated gene 5
MH	Mechanic's hands
MHC	Major histocompatibility complex
mRNA	messenger ribonucleoprotein
MSA	Myositis-specific autoantibody
NSIP	Non-specific interstitial pneumonia
NURD	Nucleosome remodelling deacetylase
NXP-2	Nuclear matrix protein-2
PBC	Primary biliary cirrhosis
pDCs	Plasmacytoid dendritic cells
PGA	Physician's global assessment
PM	Polymyositis
RANGAP	Ran GTPase activating protein
RNHRD	Royal National Hospital for Rheumatic Diseases
RP	Raynaud's phenomenon
SAE	Small ubiquitin like modifier enzyme
SLE	Systemic lupus erythematosus
SNP	Single nucleotide polymorphism
SRP	Signal recognition particle
SSc	Systemic sclerosis
SUMO	Small ubiquitin-like modifiers
TIF1- $\gamma$	Transcriptional intermediary factor 1-gamma
TNF	Tumour necrosis factor
tRNA	transfer ribonucleoprotein
UV	Ultra-violet
VAS	Visual analogue score
VCAM-1	Vascular cellular adhesion molecule-1
UIP	Usual interstitial pneumonia

## **ABSTRACT**

**Introduction:** The idiopathic inflammatory myopathies (IIM): dermatomyositis (DM) and polymyositis (PM) have been historically defined by broad clinical and pathological criteria. These conditions affect both adults and children with clinical features including muscle weakness, skin disease and internal organ involvement. Using a clinico-serological approach DM and PM can be defined into more homogeneous subsets. Myositis-specific autoantibodies (MSAs) are directed against cytoplasmic or nuclear components involved in key regulatory intra-cellular processes including protein synthesis, translocation and gene transcription. Over the last few years MSAs have been better characterised including autoantibodies directed against the aminoacyl tRNA-synthetase (ARS) enzymes, the signal recognition particle and the Mi-2 protein.

**Aim:** The overall aim of this thesis is to describe a comprehensive clinical and serological study of adult and juvenile IIM. Autoantigen targets including novel specificities were identified using protein immunoprecipitation.

**Results:** The first part of this thesis is a descriptive study on known myositis autoantibodies in adult IIM, confirming the significant association of interstitial pneumonia with anti-ARS, severe myopathy with anti-SRP, and classic DM with anti-Mi-2 serotype. In the next section, new autoantigen systems in adult IIM are described including a new anti-ARS (anti-Zo) in the anti-synthetase syndrome. Further autoantibodies directed against small ubiquitin-like modifier enzyme and a p155/140 autoantigen are major serological subsets in adult DM, the latter significantly associated with malignancy. The final section outlines a large serological study of juvenile DM (JDM) and JDM-overlap showing the frequency and clinical associations of MSAs and myositis-associated autoantibodies, including work on two new major subsets anti-p155/140 and anti-p140, which appear to define more severe disease.

**Conclusion:** The work in this thesis highlights the importance of autoimmunity in IIM and suggests a new approach where MSAs can classify patients into clinical syndromes, which predict outcomes and may as a result influence treatment strategies.

## CHAPTER 1

### INTRODUCTION

#### 1.1 Background

The idiopathic inflammatory myopathies (IIM) are a group of rare autoimmune diseases that affect both adults and children. The major subtypes of IIM are polymyositis (PM) and dermatomyositis (DM) that are heterogeneous conditions characterised by skeletal muscle weakness, biochemical or histological evidence of muscle inflammation, skin lesions and systemic organ involvement (1, 2). Morbidity outcomes relate to well-known complications, e.g. cancer-associated myositis and interstitial pneumonia. In children there is also a significant morbidity associated with vasculopathic organ involvement, soft tissue calcinosis and skin ulceration. Response to immunosuppressive therapies is variable and patients may die from disease or treatment complications, or remain very disabled. The associations with cancer and interstitial pneumonia make IIM an important autoimmune model to investigate further. The precise aetiology is unknown, mechanisms require further elucidation but the interaction between immunogenetic and environmental factors appears to play a major role (3). Like other systemic autoimmune diseases including systemic lupus erythematosus (SLE) and systemic sclerosis (SSc), IIM is characterised by the production of autoantibodies. The identification and characterisation of autoantibodies is an important cornerstone in the diagnosis of autoimmune connective tissue disease (CTD) (4, 5). Autoantibodies and corresponding target autoantigens are the subject of considerable interest, and there is increasing evidence to support a critical association between genotype, serotype and clinical phenotype in both adult and juvenile myositis (6-10).

#### 1.2 Classification and diagnosis

In 1975, *Bohan and Peter* first proposed criteria to aid the diagnosis and classification of PM and DM (1, 2). The five adapted criteria proposed were as follows:

1. Symmetrical proximal weakness of the limb or pelvic girdle and anterior neck flexors progressing over weeks to months, with or without dysphagia or respiratory muscle involvement.
2. Elevation of muscle enzymes, including creatinine kinase (CK).
3. Abnormal electromyogram (EMG) with a triad of
  - Short duration, small amplitude polyphasic motor unit potentials.

- Fibrillations, positive sharp waves, increased insertional irritability.
- Spontaneous, bizarre, high frequency repetitive discharges.
- 4. Abnormal muscle biopsy demonstrating inflammatory cell infiltrate, muscle degeneration, regeneration or necrosis.
- 5. Typical cutaneous signs of DM including
  - Heliotrope rash – violaceous erythema on upper eyelids
  - Gottron’s papules – violaceous keratotic papules on extensor aspects of fingers or elbows or knees.

Exclusion criteria: absence of other forms of myopathy e.g. inclusion body myositis, metabolic, inherited or infectious forms of myopathy.

*Bohan and Peter* suggested five subsets of myositis: PM, DM, cancer-associated myositis, juvenile myositis and myositis overlapping with CTD. *Bohan and Peter* also described that patients can develop cutaneous features of DM, which may precede the onset of myopathy (2).

- Definite DM requires all five criteria.
- Definite PM requires first four criteria (without rash).
- Probable disease requires three criteria out of first 4 criteria (for DM, one of which is rash i.e. criteria 5).
- Possible disease requires two criteria out of first 4 criteria (for DM, criteria 5 plus one other).

Overlap myositis: must fulfil criteria for their main CTD e.g. systemic sclerosis (11) or systemic lupus erythematosus (12), at least two of criteria 1 – 4, and at least one myositis-specific or associated autoantibody (see section 1.6.4).

Tanimoto *et al* published further classification criteria for PM / DM in 1995 (13).

1. Skin lesions:
  - Heliotrope rash.
  - Gottron’s papules (extensor aspect of fingers, elbows or knees).
2. Systemic inflammatory signs (elevated erythrocyte sedimentation rate, C-reactive protein, pyrexias, weight loss).
3. Non-erosive arthritis.

4. Myositis-specific autoantibodies e.g. anti-JO-1
  5. Elevated skeletal muscle enzymes.
  6. Myalgia at rest or with contraction.
  7. Proximal muscle weakness.
  8. Myopathic changes on EMG.
  9. Myositis changes on muscle biopsy.
- For a diagnosis of definite PM – at least four from item list 2 – 9.
  - For a diagnosis of definite DM – at least four from item list 2 – 9 plus at least one feature from item 1.

The IIM spectrum also includes a further subset, sporadic inclusion body myositis (IBM) and *Griggs et al* have subsequently proposed classification criteria (14). Unlike other IIM, sporadic IBM causes slowly progressive myopathy with atrophy with a distinctive pattern of muscle involvement that may be asymmetrical and affect distal rather than proximal muscle groups, and is often unresponsive to conventional immunomodulatory therapy (15). Historically, IBM cases have been seen and managed by Neurologists. Clinicians now recognise histopathology and other more specific investigations are required to exclude IBM, hereditary and other non-immune mediated myopathies when applying the *Bohan and Peter* criteria. This study has not investigated this subgroup of patients and in this thesis IIM refers purely to PM and DM cases.

### **1.3 Epidemiology**

The true frequency of PM and DM as stand-alone disorders or in association with other CTD is unknown. Reports of incidence and prevalence are limited based on different study populations. Estimates based on the diagnostic criteria proposed by *Bohan and Peter* (1, 2), may not be accurate due to difficulties distinguishing PM and IBM. Reports show an annual incidence of 2-7 / million population (16-18). The mean age of onset is 50 years affecting women more than men. Juvenile dermatomyositis (JDM) is the most common IIM of children and the reported incidence is 0.8-4.1 per million children per year (18-20). In contrast to adults, juvenile PM is rare (21), whereas JDM overlap with other connective tissue disease in particular SSc and SLE is more common.

## **1.4 Clinical features**

IIM is characterised by a spectrum of clinical features including skeletal muscle weakness, skin lesions and systemic organ involvement, in particular interstitial pneumonia and a risk of cancer in adults. Children with JDM share some clinical features with adult patients in terms of muscle disease and characteristic skin lesions. Certain clinical features including skin ulceration, calcinosis and gastrointestinal involvement are more common in children and have been proposed as predictors of severe disease course in JDM (21-25). In contrast to adults with DM, both interstitial pneumonia and cancer-associated myositis are very rare in JDM. Delays in diagnosis, in particular recognising active muscle or lung disease and poor response to treatment mean that patients are often left disabled from irreversible damage.

### **1.4.1 Myositis**

IIM can present with progressive symmetrical muscle weakness that usually develops over a few weeks to months. The proximal muscle groups such as the shoulder girdle, pelvic girdle, thigh muscles and neck flexors are typically involved. In severe cases, other muscle groups can be involved including chest wall weakness, which can compromise respiratory function. At onset, patients rarely complain of muscle pain but experience difficulties with activities of daily living, for example, rising from a chair, walking up the stairs and raising their arms. Depending on the severity of the case and the response to treatment muscle atrophy can develop that leads to severe irreversible weakness.

### **1.4.2 Cutaneous manifestations**

Classic DM skin lesions are generally present at the time of diagnosis and in some cases may precede the development of muscle inflammation or other organ involvement (see Section 1.5.3) (26). Gottron's sign is an erythematous, scaly plaque-like lesion usually found over the dorsal surface of the metacarpophalangeal or proximal interphalangeal joints and the extensor surfaces of the knees or elbows. Heliotrope rash is a purple violaceous eruption of the upper eyelids, often with cutaneous oedema. More extensive skin involvement includes the V-sign and Shawl-sign rash over the anterior triangle of the neck, and the back of the neck and shoulders respectively. Erythroderma can develop on the face (malar distribution or forehead). All these rashes are diffuse, flat and erythematous and may be photosensitive. The skin lesions can have a poikilodermatous appearance with a combination of atrophy, pigment changes and telangiectasia. Periungual erythema



with abnormal dilated nailfold capillaries is another feature. Mechanic's hands are a characteristic cutaneous manifestation that is pathognomonic of the clinical and serological subset termed the anti-synthetase syndrome (see Section 1.5.1). The description refers to dry, cracked or fissured skin with hyperkeratosis over the lateral aspects of the inter-digits especially the index and middle fingers. Certain features secondary to persistent cutaneous vasculitis are more common in JDM, in particular generalised oedema, skin ulceration and soft tissue calcification (calcinosis) (21-25).

### **1.4.3 Interstitial pneumonia**

Diffuse parenchymal lung disease (DPLD) is a major cause of morbidity and mortality in IIM (27). Primary lung involvement or ventilatory failure secondary to muscle weakness is common, more so in adult IIM (28-30). Interstitial pneumonia is the most frequent sub-type of DPLD in IIM, with studies showing an overall frequency of between 20-60% (29, 31). Interstitial pneumonia (also termed interstitial lung disease) are a heterogeneous group of non-malignant and non-infectious lung disorders that comprise of a number of clinico-pathological entities that are sufficiently different to be designated as separate diseases. The characteristic hallmark is that of varying patterns of inflammation and fibrosis affecting the lung parenchyma.

In 2002 the American Thoracic Society and the European Respiratory Society (ATS/ERS) published the consensus classification for interstitial pneumonia that is equally applicable in CTD (32). Based on radiological and histopathological patterns, interstitial pneumonia can be further subdivided into several subsets including usual interstitial pneumonia (UIP), non-specific interstitial pneumonia (NSIP), organising pneumonia (OP) and acute interstitial pneumonia (AIP), which are recognised in CTD (32, 33). The histological hallmark of UIP is the presence of fibroblastic foci with a subpleural, basal and peripheral distribution (32, 34). The most important differential pattern of UIP, particularly in CTD is NSIP, which has a similar subpleural and basal distribution. *Katzenstein et al* first described NSIP characterised by cellular and fibrotic changes with the absence of dense fibrosis or fibroblastic foci (35). With the advent of high-resolution computerised tomography (HRCT), interstitial pneumonia patterns in CTD can be categorised radiologically with good histological correlation (36, 37).

NSIP has been reported to be the most frequent subtype on both HRCT and lung biopsy in IIM (38). AIP, COP and UIP have also been reported and identified as

patterns responsible for acute or rapidly progressive interstitial pneumonia with AIP and UIP associated with the worst prognosis (28-31, 33, 39). It is well recognised that interstitial lung involvement may be the first or predominant organ manifestation preceding muscle or skin disease, particularly in the anti-synthetase syndrome (see Section 1.5.1) and in patients with clinically-amyopathic DM (CADM) (see Section 1.5.3) (29, 31, 40-42).

#### **1.4.4 Arthritis and overlap features**

Non-erosive inflammatory arthritis is a frequent manifestation of IIM, particularly in anti-synthetase syndrome and overlap conditions such as SSc, mixed connective tissue disease and SLE. Raynaud's phenomenon characterised by vasospasm and structural changes of the digital microvasculature is a common feature. Joint disease with contractures and lipoatrophy of soft tissues is more frequent in JDM particularly in patients with overlap scleroderma. General features include fatigue, weight loss and fever.

#### **1.4.5 Other organ manifestations**

Other organs can be involved including the striated muscles of oropharyngeal and upper oesophagus causing dysphagia, and gastro-intestinal tract vasculopathy leading to ischaemic ulceration with pain and bleeding, the latter a more common complication in JDM (43). Cardiac involvement including myocarditis has been reported with subsequent conduction defects and ventricular failure (44, 45). Although rare, cardiac disease confers a high risk for mortality in patients with IIM (27, 44).

### **1.5 Specific clinical subsets**

#### **1.5.1 Anti-synthetase syndrome**

The anti-aminoacyl tRNA synthetase (anti-ARS) autoantibodies (see Section 1.6.2) define the anti-synthetase syndrome recognised as a spectrum of myositis, interstitial pneumonia, non-erosive arthritis, fever, Raynaud's phenomenon and mechanic's hands (46). Patients may have different clinical features within the syndrome, but overall there is a significant association with interstitial pneumonia and studies have described the frequency to be as high as 95% (28-30, 47, 48).

### **1.5.2 Cancer-associated myositis**

The association between cancer and adult DM is well recognised particularly in the older age groups (49-54). Because IIM can have a chance association with malignancy, cancer-associated myositis is defined by a temporal association of less than three years between cancer and DM onset (50, 51). The association with PM is uncertain with some studies reporting an increased relative risk but some not (53, 55, 56). Hill *et al* conducted a pooled analysis of published national data from Sweden, Denmark, and Finland on all patients with DM and PM (over the age of 15 years). The study calculated standardised incidence ratios (SIR) for individual cancer sites for dermatomyositis and polymyositis separately, using national cancer rates by country, sex, age, and date. The frequency of cancer in the DM cohort was 32% (58% of which developed cancer after the diagnosis of DM), with a SIR of 3.0. The most common cancer observed was ovarian followed by lung, pancreatic, non-Hodgkin lymphoma stomach and colorectal. 15% of PM cases were associated with cancer (70% developed after the diagnosis of PM), with a SIR of 1.3. PM was associated with a higher risk of non-Hodgkin lymphoma, then lung and bladder malignancy. In both DM and PM, the risk of malignancy was highest at the time of myositis diagnosis (within the first year) with a markedly reduced relative risk over time. For DM the SIR of cancer by year after diagnosis of myositis was 13.5 (0-1 year), 2.5 (2-5 years) and 1.4 (>5 years). In the PM group the SIR of cancer by year after diagnosis of myositis was 2.6 (0-1 year), 1.5 (2-5 years) and 0.9 (>5 years). In the DM cohort the number of cancers recorded in patients under the age of 44 years was small, and there was no increase in risk in PM patients (53). In contrast to adults, JDM appears not to be associated with malignancy (53, 57). Historically, cancer-associated myositis has not been associated with specific myositis autoantibodies, but recent reports including work described in this thesis have described a novel autoantibody marker in this clinical subset (see section 1.6.2).

### **1.5.3 Clinically-amyopathic dermatomyositis (CADM)**

Patients may only develop DM skin changes with no clinical evidence of myopathy, termed DM sine myositis or amyopathic DM (ADM) (40). Clinically-amyopathic DM (CADM) is a designation for patients who either have ADM or those patients who have DM skin lesions, no muscle weakness but evidence of sub-clinical myopathy on further testing (muscle enzymes, EMG, imaging or biopsy), termed hypomyopathic DM.

## 1.6 Treatment

The treatment of both PM and DM is often empirical and in general is influenced by clinical experience rather than robust clinical data. There remains a paucity of standardised clinical trials due to the relative rarity of PM and DM. In addition, perhaps due to historical classification tools including the *Bohan and Peter* criteria, IIM is a heterogeneous condition, which makes designing trials and interpreting treatment outcomes in different disease subtypes difficult. For example, optimal treatment strategies for ASS with lung disease may be different to other subgroups, CADM or immune-mediated necrotising myopathy. Moreover, the absence of standardised disease activity and outcome measures has made the assessment of treatment response difficult. Classifying patients into more homogenous subsets based on serotype / clinical phenotype, and the advent of validated myositis disease activity tools (see section 1.7) will facilitate the design of clinical trials, which can capture clinically significant changes.

Based on personal clinical experience and limited trial data there are several treatments that are effective treating patients with IIM.

### ***Glucocorticoids***

Prednisolone is recognised as the first line treatment for PM or DM including those with internal organ complications e.g. ASS with interstitial lung involvement. This is primarily based on clinical experience rather than trial data. In the majority of cases, the initial dose is 0.5-1mg/kg daily, and depending on clinical response the dose is weaned over a number of months. Patients may remain on low dose prednisolone for a number of years, particularly those with ASS and ILD, which appears to be very steroid responsive but also steroid dependant with disease flares on steroid withdrawal (48) (personal observation). In severe cases of myositis with internal organ involvement (including those with gastrointestinal involvement e.g. vasculopathy or pharyngeal dysfunction), patients may require induction with intravenous methylprednisolone pulses.

### ***Specific immunomodulatory therapy***

In patients who respond to steroids, the goal is to induce disease control and thus remission as quickly as possible at the same as minimising steroid requirements. In mild to moderate cases, for example those with purely muscle and / or skin disease, this may be achieved with either Azathioprine or Methotrexate (58-61). An alternative to Azathioprine and Methotrexate is Ciclosporin, the latter two agents

have also been compared in a small randomised controlled trial, with no significant difference observed, although appeared effective (62). However, the side effect of profile may be higher with Ciclosporin, which often limits its use in clinical practice. In moderate to severe cases, induction therapy in combination with pulsed / oral prednisolone may be required. The efficacy of intravenous immunoglobulin (IV IG) has been demonstrated, especially in DM, in a randomised controlled trial (63). In DM patients who response to standard regimes with prednisolone and Azathioprine / Methotrexate is inadequate or in those patients with severe disease at presentation (not complicated by interstitial pneumonia), IV IG is the preferred agent of choice. Repeated infusions may be required to achieve disease control followed by maintenance therapy as outlined above. In severe cases, including those with significant interstitial pneumonia and / or severe necrotising myopathy, based on personal clinical experience, IV Cyclophosphamide (CYC) may be a useful adjuvant therapy. In general, induction with 4-8 pulsed infusions given every 3-4 weeks is an optimal strategy. If this treatment fails or is contra-indicated, the novel biologic agent Rituximab (monoclonal antibody against CD20-positive B cells) appears to be an attractive treatment option, especially in myositis-autoantibody-positive refractory patients e.g. ASS patients with interstitial pneumonia and DM (64). A multi-centre / international randomised placebo controlled has recently completed recruitment (<http://www.edc.gsph.pitt.edu/rimstudy/index.html>). There is now preliminary data for newer agents in moderate disease or as maintenance therapy following induction with IV IG, IV CYC or Rituximab. Mycophenolate Mofetil inhibits the *de novo* pathway of nucleotide synthesis leading to an anti-proliferative affect directed against T and B lymphocytes. It appears to be a useful agent particularly in those patients where Azathioprine or Methotrexate is ineffective or poorly tolerated. There is also preliminary data that suggests Mycophenolate may be beneficial in refractory IIM cases associated with interstitial pneumonia (65-67). A further option is Tacrolimus, a calcineurin inhibitor that blocks T cell signal transduction and IL-2 signalling. This drug can be effective in some difficult-to-treat cases of polymyositis / ASS, especially in patients with interstitial lung disease (68, 69).

### **1.7 Assessment Tools**

In a clinical setting, proximal myopathy is often assessed by physician tested medical research council (MRC) scale for muscle strength. Manual muscle testing, however, has a number of limitations. One limitation is that the MRC scale is an ordinal scale with disproportional distances between grades. Another limitation of the MRC scale is that the scoring depends on the judgment of the examiner. Finally,

with the 6-point ordinal MRC scale, it is difficult to identify relatively small but clinically relevant changes in muscle strength. Alternative more objective tests to assess proximal muscle function have been devised, including two isotonic tests (1 kg arm lift for upper limb function and a 30 second chair stand test for lower limb function). These tests exhibit excellent test-retest reliability, demonstrate construct validity and are responsive to changes in disease activity in IIM (70). New tools have been devised to standardise the conduct and reporting on IIM clinical trials. Two tools, known as the myositis intention to treat index (MITAX) and the myositis disease activity assessment visual analogue scale (MYOACT), have been developed to measure activity in patients with IIM ([www.niehs.nih.gov/research/resources/collab/imacs/docs/activity/MDAAT\\_Scoring\\_2009.pdf](http://www.niehs.nih.gov/research/resources/collab/imacs/docs/activity/MDAAT_Scoring_2009.pdf)). In addition, the myositis damage index (MDI) has been devised to assess the extent and severity of damage developing in different organs and systems. These measures have been reviewed by the International Myositis Assessment and Clinical Studies (IMACS) group and have been found to have good face validity (71). A recent study has demonstrated that the myositis disease activity assessment tool (MITAX and MYOACT) has good interrater reliability and validity in clinical practice (72). In addition to these tools, specific disease assessment scales have also been designed in juvenile IIM, which are used in routine clinical practice, as well as research studies (see section 5.1.2 and appendix 5.5).

## **1.8 Pathogenesis and aetiology**

It is becoming increasingly clear that IIM is a useful model to study the relationship between immunogenetic profiles, autoimmune targets and clinical phenotype in CTD. The development of IIM appears to be secondary to the combination of genetic factors and exposure to environmental agents that may initiate specific autoimmune responses in certain individuals.

### **1.8.1 Pathogenesis**

#### ***Classic descriptions***

Cellular and humoral autoimmune mechanisms that target skeletal muscle, the lungs, the skin, and associated microvasculature are implicated in the pathogenesis of IIM. In broad terms, in DM, immune responses are said to primarily target the microvascular endothelium leading to activation of the complement cascade, upregulation of cytokines and chemokines, and thus endothelial damage. Inflammatory lesions with B and CD4 positive T cells are predominately found in perivascular and perimysial regions with perifascicular atrophy. Endothelial

hyperplasia and fibrin thrombi are seen in both endomysial and dermal capillaries (17, 73, 74). Cytokines are released, which leads to upregulation of vascular-cell adhesion protein-1 (VCAM-1) and intercellular adhesion molecule (ICAM-1), which promotes endothelial damage. Historically in PM, myocytotoxicity is said to be mediated by cellular responses where clonally expanded CD8 positive T cells are primed against muscle fibres expressing MHC class I antigens leading to an inflammatory cell infiltrate that invades muscle fascicles (17, 75). Previous descriptions have suggested B cells are sparse within the inflammatory cell infiltrate of PM specimens (76).

### ***Shared cellular and humoral mechanisms***

It is now becoming increasingly clear that both T and B cell responses occur in both subtypes of IIM. Both CD4 and CD8 T cells have been demonstrated in PM and DM muscle tissue (77). In addition, cytokines, chemokines and their receptors, are strongly expressed by endothelial and inflammatory cells in both subtypes. CD4+ cells previously characterised as T cells have been identified as plasmacytoid dendritic cells (pDCs). Large-scale gene expression studies have confirmed a type 1 interferon (IFN) signature with abundant pDCs in DM muscle and skin (78, 79). More recently, type 1 IFN  $\alpha/\beta$  producing pDCs have also identified in PM muscle (80), highlighting the potential role of this cytokine system in the pathogenesis of IIM. Furthermore, antigen-driven B cell humoral responses have now been demonstrated within muscle in PM, as well as DM. A previous study has shown that immunoglobulin transcripts are prominent in PM muscle, with a relative abundance of CD138+ plasma cells (81). Further work has confirmed local B cell maturation occurs within myositis muscle in both subtypes of IIM; B cells and corresponding plasma cells were clonally expanded, had class-switched and undergone mutation (82). This suggests antigen-driven responses may play a crucial role in disease pathogenesis. The role for B-cell autoimmunity in PM and DM is further emphasised by the production of autoantibodies against specific intra-cellular protein targets, including the aminoacyl-tRNA-synthetases, signal recognition particle and Mi-2.

### ***Non-immune mechanisms***

It has been suggested that non-immune mechanisms may play a role in disease pathogenesis. The rationale being some patients can have minimal inflammatory cell infiltrate but marked structural changes on muscle biopsy with significant weakness. In a mouse model, over-expression of class I MHC molecules has been shown to mediate skeletal muscle fibre damage, even in the absence of T and B

cells (83). Similarly, in human myositis, induction of class I MHC in muscle fibres occurs before mononuclear cell infiltration (84, 85). Recent data suggests continuous upregulation of MHC class I on muscle fibers leads to an endoplasmic reticulum (ER) stress response. A potential mechanism has been proposed whereby ER overload leads to activation of nuclear factor kappa-B pathway combined with the ER unfolded protein response, which in turn promotes pro-inflammatory cytokine activity, cell death and muscle fibre necrosis (86).

### **1.8.2 Immunogenetics**

Evidence for a genetic basis for IIM has largely accumulated through candidate gene studies, although a familial predisposition is suggested from case reports of familial clustering (87). Candidate gene studies have confirmed that HLA-DRB1\*0301 and HLA-DQA1\*0501 are definite risk factors for the development of Caucasian IIM (9, 87-89). These alleles form part of the conserved, ancestral Caucasian haplotype, A1-B8-Cw7-DRB1\*0301-DQA1\*0501. Recent studies have confirmed IIM in different ethnic groups are associated with different HLA-DR and DQ alleles (8, 89-91).

Major histocompatibility complex (MHC) differences exist between the traditional clinical PM and DM sub-types, where HLA-DRB1\*07 confers risk for DM but is protective for PM (9). The 8.1 ancestral MHC haplotype, (HLA-B\*08-DRB1\*03-DQA1\*05-DQB1\*02) is associated with adult and juvenile IIM (9, 10), where the strongest signal appears to come from the HLA class I region (89, 92). Furthermore, TNF-308A single nucleotide polymorphisms (SNP) are associated with IIM, but only due to strong linkage disequilibrium with HLA-B\*08, thus forming part of an extended 8.1 haplotype (92). Clinical phenotype is more strongly associated with myositis autoantibodies than with traditional clinical sub-type in adult Caucasian IIM (9, 89). HLA class II haplotype (DRB1-DQA1-DQB1) is in turn associated with antibody status. For example in adults, the DRB1\*07-DQA1\*02-DQB1\*02 haplotype is strongly associated with anti-Mi-2, whereas the 8.1 haplotype is strongly associated with anti-Jo-1 autoantibodies (see also Section 1.6.2) (9).

### **1.8.3 Environmental factors**

Environmental triggers may also influence the development of myositis. Evidence implicating potential environmental factors includes the observation that seasonal patterns are associated with the development of specific IIM-serological syndromes e.g. anti-Jo-1 positive patients have been shown to develop disease in the Spring



(93, 94). In a recent study, seasonal birth patterns in myositis subgroups suggested an aetiological role of early environmental exposures (95). In particular, birth distributions appeared to have a stronger seasonal association in JDM, suggesting a potential role for specific perinatal exposure.

There is evidence to suggest an aetiological role of ultra-violet (UV) light in the pathogenesis of autoimmune CTD. From a clinical perspective SLE and DM are associated with photosensitive rashes, and disease flares can be induced by sunlight exposure. Studies have demonstrated that UV-B light induces apoptosis of keratinocytes, which leads to the redistribution of certain intracellular proteins that are classically recognised by SLE-associated autoantibodies (96-98). In particular, autoantigens were shown to be clustered into two distinct populations, either in surface blebs or larger surface bodies on apoptotic cells (96). Apoptotic blebs contained cytoplasmic components including fragmented endoplasmic reticulum and the Ro52 protein. Apoptotic bodies contained nuclear associated autoantigens, nucleosomal DNA, Ro60, La, and the small nuclear ribonucleoproteins. These findings implicate apoptotic cells as an important potential source of concentrated or modified autoantigens that may drive autoantibody production and skin disease in SLE (97). Similar to SLE, there is evidence to suggest an aetiological role of UV-light in the pathogenesis of DM. Firstly, the incidence of DM cases has been shown to be higher in countries with a latitudinal gradient towards the equator (99). Secondly, surface UV-radiation intensity has been shown to be associated with the relative frequency of DM cases, especially those with anti-Mi-2 autoantibodies (classically associated with hallmark cutaneous DM lesions, see section 1.6.4.1). Collectively, this data provides insight into the possible aetiological mechanisms of the development of DM.

#### **1.8.4 Autoantibodies and targeted autoantigens**

The identification and characterisation of autoantibodies is an important cornerstone in the diagnosis of autoimmune CTD (4, 100, 101). Autoantibodies target nuclear or cytoplasmic intra-cellular components and specific profiles are associated with clinical phenotypes within the CTD disease spectrum. Autoimmunity is characterised by the loss of tolerance to self-proteins or autoantigens. In recent years there have been several attractive paradigms proposed to explain why certain seemingly ubiquitously expressed proteins are selectively targeted by autoantibodies. Autoantigens appear to be critical partners in driving autoreactivity, unified by their susceptibility to cleavage by certain apoptotic substrates. Studies have

demonstrated the protease granzyme B, an enzyme released by cytotoxic T cells, generates a unique form of apoptosis that leads to the cleavage of certain intracellular proteins into novel fragments. These fragments become clustered and may trigger autoreactive responses. This mechanism has been strongly implicated in the pathogenesis of several autoimmune diseases including SLE, SSc and IIM (96, 97, 102-106). Furthermore, this apoptotic trigger is likely to occur in specific pro-immune microenvironments and there is now emerging evidence to suggest that specific autoantigens are upregulated in different cell types and lesional tissues, particularly in IIM (107, 108).

#### **1.8.4.1 Autoantibodies in myositis**

Previous studies have shown that autoantibodies are detected in approximately 50-60% of IIM patients depending on the detection methods used (46, 109, 110). The standard techniques have included immunofluorescence on human-epithelial cell lines (HEp-2 cells), generic or antigen-specific ELISA tests, immunodiffusion and counter-immunoelectrophoresis. More specific techniques including immunoblotting and in particular protein radioimmunoprecipitation (IPP) have the advantage of being able to detect novel autoantigen targets. The major advantage of IPP is that it identifies those autoantibodies that target conformational epitopes. In addition, novel targets are amenable to identification using a proteomic approach. The identification and characterisation of novel autoantibody subsets in IIM in this study are based on the method of IPP.

Autoantibodies in IIM can be divided broadly into two groups, myositis-specific autoantibodies (MSAs) or myositis-associated autoantibodies (MAAs). MAAs are found in myositis overlap syndromes, in particular SSc, MCTD and SLE. In general, MSAs are unique to PM and DM, and the previously described specificities are outlined below. See Tables 1 and 2, and Figure 1.

### ***Anti-synthetase autoantibodies***

Autoantibodies against the cytoplasmic aminoacyl-tRNA synthetase enzymes (ARS) are the most frequently detected in adult patients with myositis. The first to be described and most common autoantigenic target is Jo-1 (histidyl tRNA-synthetase) (111), which is found in approximately 20% of IIM patients (112, 113). Autoantibodies to six other ARS molecules; PL-7 (threonyl), PL-12 (alanyl), EJ (glycyl), OJ (isoleucyl), KS (asparaginy) and Ha (tyrosyl) have been described collectively in approximately 20% of patients with IIM (frequency of patients with each non-Jo-1 anti-ARS is between 1-5% (114-119). A further anti-ARS termed anti-Zo (phenylalanyl) has been discovered by our group and will be described in Chapter 4 (120). The ARS are a distinct group of enzymes that catalyse the binding of specific amino acids to their cognate tRNA for incorporation into growing polypeptide chains. The anti-ARS autoantibodies define the anti-synthetase syndrome, which has been described in Section 1.5.1.

### ***Anti-SRP autoantibodies***

Patients with autoantibodies to the signal recognition particle (SRP) appear to form a further distinct clinical and histopathological subset. *Reeves et al* were the first to report SRP as an autoantigen target in myositis (121). The SRP autoantigen is a ribonucleoprotein complex (proteins-7SLRNA) with the 54 kDa and 72 kDa polypeptides most frequently targeted. The cytoplasmic SRP protein recognises secretory or membrane bound proteins and regulates protein translocation through the endoplasmic reticulum. Anti-SRP autoantibodies are found in approximately 5% of patients with IIM (122, 123). Patients with anti-SRP can present with acute onset severe myopathy with significant muscle enzyme elevation and systemic features including dysphagia that can be refractory to standard treatments (123, 124).

### ***Anti-Mi-2 autoantibodies***

Anti-Mi-2 is classically detected in adult and juvenile patients with DM clinical features. Anti-Mi-2 is detected in between 5-20% of patients with DM (6, 10, 125-128). Mi-2, a nuclear helicase protein forms part of the nucleosome remodelling deacetylase (NuRD) complex, which plays a role in gene transcription (129). This autoantibody specificity is described in patients with hallmark cutaneous DM lesions including Gottron's papules, heliotrope rash, cuticular overgrowth and the V-sign / Shawl-sign. Moreover, patients may have milder muscle involvement with a lower risk of interstitial pneumonia and respond well to therapy (6, 125-127).

***Anti-CADM-140 autoantibodies***

A novel autoantibody specificity termed anti-CADM-140 in a Japanese cohort associated with CADM and rapidly progressive interstitial pneumonia (130). This autoantibody specificity was not seen in classic DM, PM or any other CTD cases. Using a series of molecular techniques the CADM-140 autoantigen has recently been identified as the cytoplasmic protein melanoma-differentiation associated gene 5 (MDA5) (131). MDA5 is involved in innate immune responses against viral infections (132).

**Table 1: Myositis-associated autoantibodies, autoantigen targets and clinical features**

<b>Autoantibodies</b>	<b>Autoantigen and function</b>	<b>Clinical Phenotype</b>
Anti-U1-RNP	snRNP ribonucleoprotein complex Splicing mRNA	Mixed connective tissue disease – overlap features including myositis, arthritis, sclerodacty, Raynaud’s, diffuse parenchymal lung disease, pulmonary hypertension
Anti-Ro52/60	Ro ribonucleoprotein complex (proteins hYRNA)  Binding to DNA and transcription factors	SLE, Sjogren’s, overlap myositis, reported in association with anti-ARS autoantibodies
Anti-PM-Scl	Nucleolar multi-protein Biogenesis of ribosomes	Myositis-scleroderma overlap
Anti-U3-RNP	Nucleolar fibrillarin ribonucleoprotein Pre-ribosomal RNA processing	Myositis-scleroderma overlap
Anti-Ku	Heterodimer Ku-p350 complex DNA repair and phosphorylation of transcription factors	Myositis-scleroderma overlap

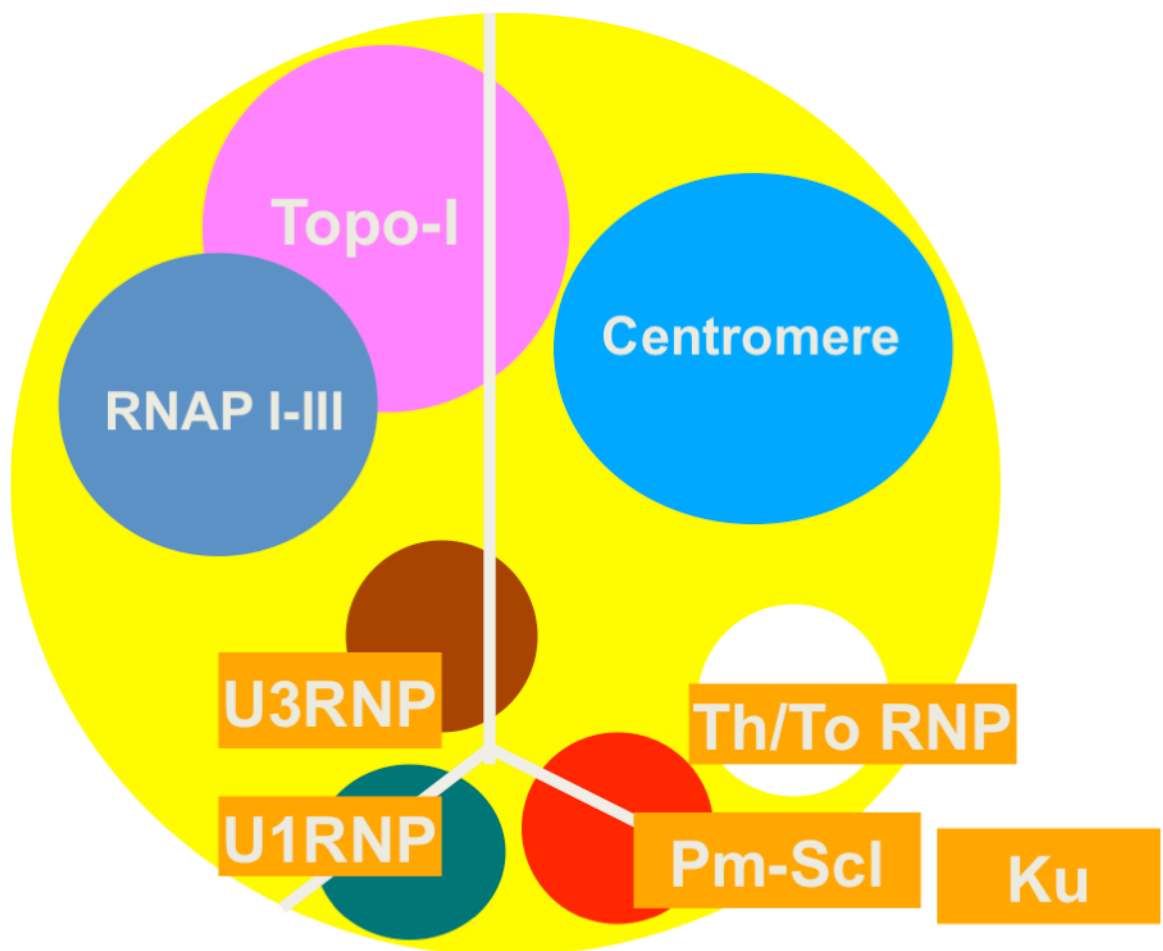
**Table 2: Myositis-specific autoantibodies, autoantigens and clinical features**

<b>Autoantibodies</b>	<b>Autoantigen and function</b>	<b>Clinical Phenotype</b>
Anti-ARS	ARS – intracytoplasmic protein synthesis (binding of specific amino acid to its cognate tRNA for incorporation into polypeptide chains)	Anti-synthetase syndrome
Anti-Jo-1	Histidyl	Myositis, mechanic's hands, Gottron's papules, arthritis, fever, Raynaud's phenomenon, high frequency of interstitial pneumonia
Anti-PL-7	Threonyl	
Anti-PL-12	Alanyl	
Anti-EJ	Glycyl	
Anti-OJ	Isoleucyl	
Anti-KS	Asparaginy	
Anti-Ha	Tyrosyl	
Anti-SRP	SRP – intracytoplasmic protein translocation from ribosome into endoplasmic reticulum (6 polypeptides and ribonucleoprotein 7SLRNA)	Acute onset necrotizing myopathy
Anti-Mi-2	Helicase nuclear protein - transcription, remodelling of nucleosomes	Hallmark cutaneous DM, milder muscle disease
Anti-CADM-140	Intracytoplasmic MDA5 – innate immune responses against viral infections	CADM Rapidly progressive interstitial pneumonia

ARS, aminoacyl-tRNA synthetases; SRP, signal recognition particle; NuRD, nucleosome remodelling histone deacetylase; MDA5, melanoma-differentiation associated gene 5; DM, dermatomyositis; CADM, clinically-amyopathic dermatomyositis.

**Figure 1: Autoantibodies in myositis and systemic sclerosis**

**1A: Systemic sclerosis and myositis overlap**



**Systemic sclerosis specific:**

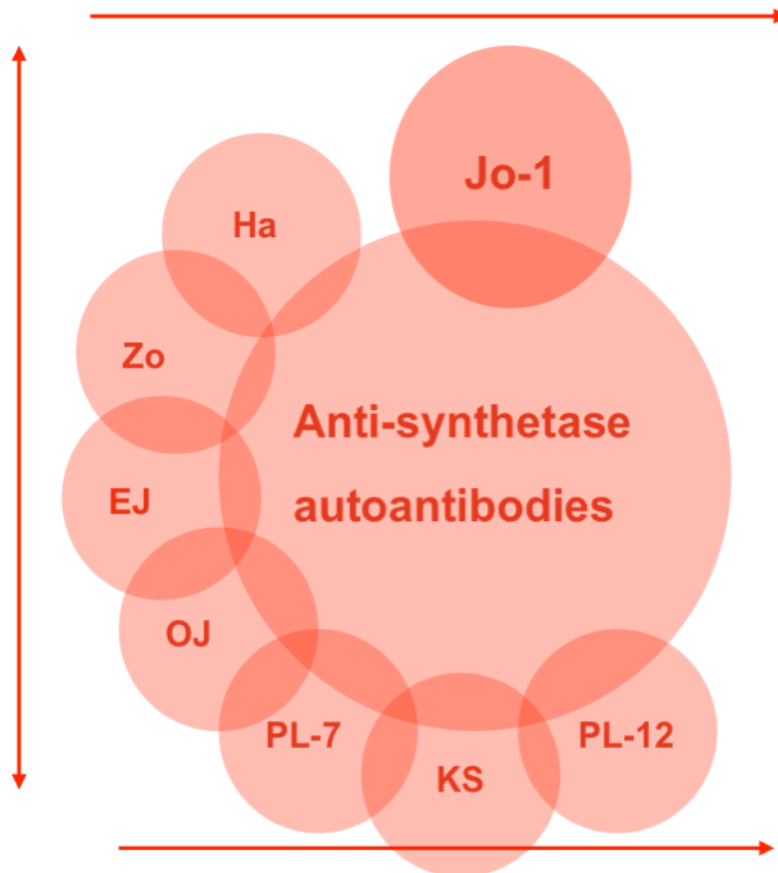
- Topo-I: Topoisomerase
- RNAP I-III: RNA polymerase I-III
- Centromere
- Th/To RNP

**Myositis-associated (overlap with systemic sclerosis):**

- U1RNP
- U3RNP
- PM-Scl
- Ku

**Figure 1**

**1B: Anti-synthetase syndrome**



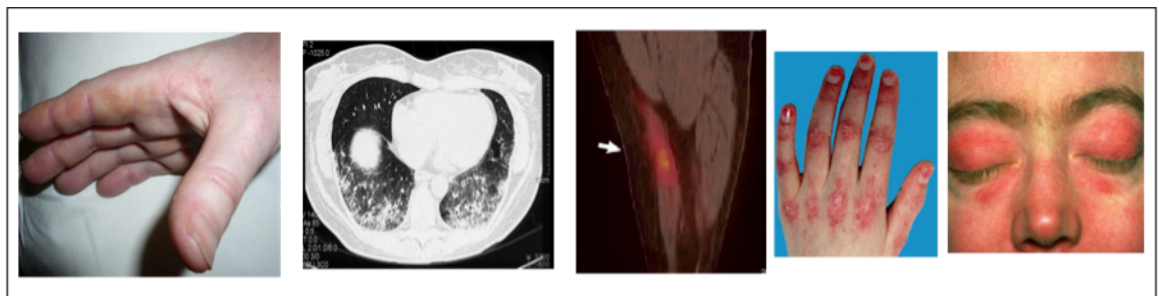
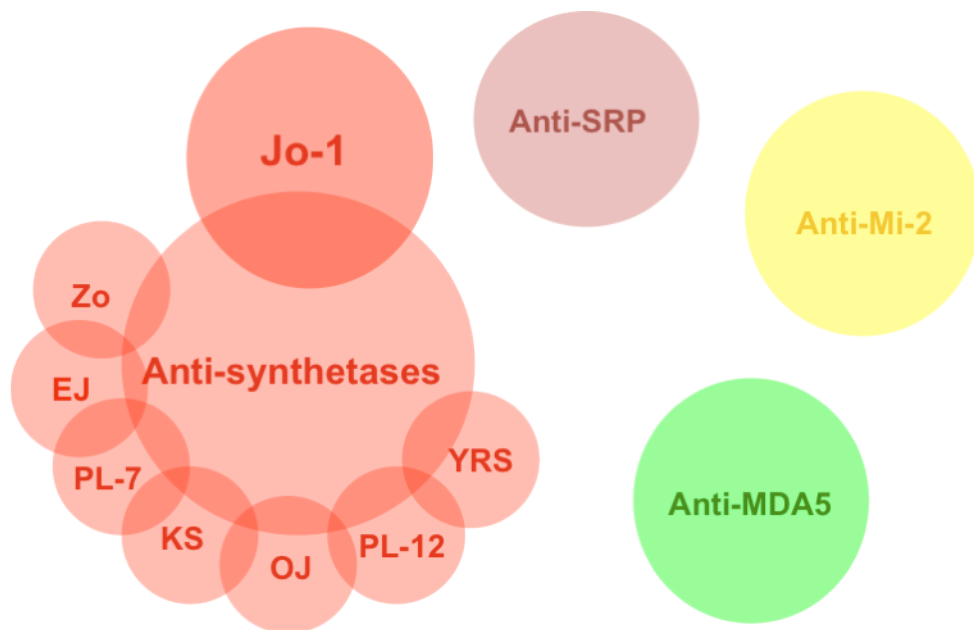
**Anti-synthetase syndrome**

- Myositis
- Raynaud's phenomenon
- Arthritis
- Mechanic's hands
- Gottron's lesions
- Fever
- Interstitial pneumonia



**Figure 1**

**1C: Myositis-specific autoantibodies**



- Anti-synthetase – anti-synthetase syndrome
- Anti-SRP – acute necrotising myopathy
- Anti-Mi-2 – classic DM with hallmark skin disease
- Anti-MDA5 – clinically-amyopathic DM with significant lung involvement

#### **1.8.4.2 Autoantigens and disease mechanisms**

Identification of autoimmune targets in IIM has led to further insights into pathogenesis. Perhaps the most striking feature is that groups of autoantigens with analogous cellular functions are associated with similar clinical subsets. For example, the target molecules in the anti-ARS autoantibody phenotype (anti-synthetase syndrome) and the anti-SRP autoantibody phenotype (severe necrotizing myopathy) are found in the cytoplasm with distinct functions involved in protein synthesis and translation. In contrast, the other myositis autoantigen targets are all nuclear proteins involved in gene transcription and DNA processing. Recent work has proposed the hypothesis that the dysregulated autoimmune response in myositis syndromes is driven by specific antigens in distinct tissue microenvironments. Two studies have demonstrated that certain autoantigens are enriched in lesional tissue involved in IIM (107, 108). Several myositis-specific and associated autoantigens were shown to be upregulated in myositis muscle in comparison to normal muscle especially in regenerating muscle cells. Of further interest was the observation that Mi-2 was preferentially expressed in DM muscle rather than PM muscle. The theory that distinct microenvironments may shape disease expression was emphasised with the finding that a novel conformation of Jo-1 following cleavage by granzyme B is enriched in the alveolar-epithelial layer of the lung in comparison to other tissues including muscle (108). This has led to the suggestion that the initiating target tissue for the autoimmune response in the anti-Jo-1 syndrome is the lung with secondary attack to muscle. Further evidence implicating autoantigen driven responses has been suggested by an antigen-induced model of IIM following immunisation of congenic mice with murine Jo-1 protein (133). Antibody responses were species-specific and following immunisation mice developed the ASS clinical phenotype with muscle and lung inflammation. The potential role of other autoantigens driving autoimmunity in IIM has been highlighted by work on the chromatin remodeler Mi-2. Using a conditional transgenic mouse model, a major function of the Mi-2 protein has been elucidated (134). In particular, Mi-2 is essential for development and repair of the basal epidermis, this observation may give further insight into the pathogenic mechanisms of the anti-Mi-2 autoantibody clinical subset, of which the predominant clinical manifestation is skin disease. Further evidence to suggest a central pathogenic role of the autoantigen in IIM comes from the demonstration that certain autoantigenic ARS proteins, histidyl (Jo-1) and asparaginyl (KS) can induce leukocyte migration. In contrast, non-antigenic ARS (aspartyl-tRNA and lysyl-tRNA synthetases) were not chemotactic. The study proposed that autoantigenic ARS are over-expressed in damaged muscle

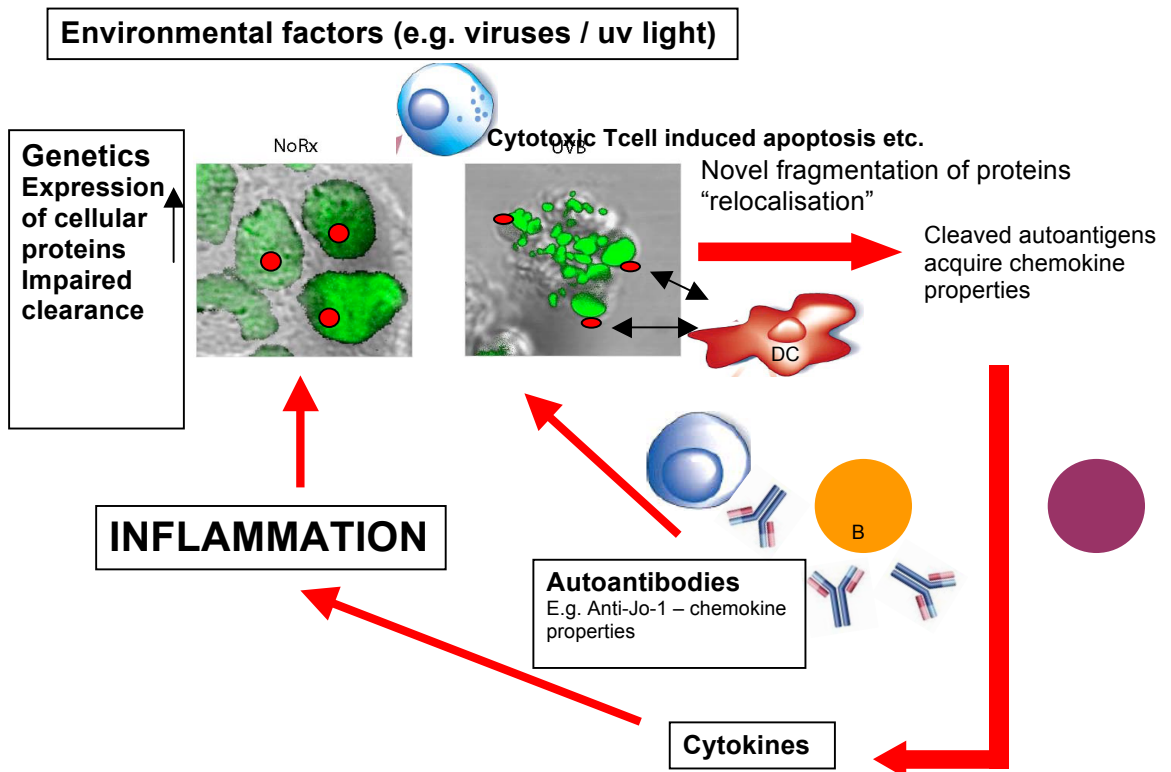
cells; their pro-inflammatory properties promote the immune response and thus the development of myositis (135). See Section 1.6.5, Figure 2.

#### **1.8.4.3 MSAs and histopathological associations**

There is also evidence to suggest that there are histopathological associations with certain MSAs, which highlights the overlap between the broad IIM subtypes. For example, patients with ASS associated myositis have distinct histopathological features with prominent perimysial inflammation, connective tissue fragmentation and perifascicular myopathic changes, in contrast to endomysial inflammation (136). Also, typical histopathological changes have been described in muscle biopsies from patients with anti-SRP autoantibodies; sparse inflammatory cell infiltrate, prominent muscle fibre necrosis and endomysial fibrosis (123, 124).

### 1.8.5 A proposed model for autoimmunity in IIM

Figure 2: Initiation of autoimmunity in IIM



Based on their studies, *Casciola-Rosen and Rosen* have proposed a hypothesis that the dysregulated autoimmune response in myositis syndromes is driven by novel patterns of antigen expression or conformation in distinct tissue microenvironments (107, 108, 137, 138). It is likely that a specific stimulus, for example tissue injury secondary to an infective or a toxic agent leads to a perturbed pro-immune setting in a genetically susceptible individual. In IIM, there may be three or four key initiating microenvironments i.e. muscle, lung or perhaps even skin or tumour tissue. As a result, altered antigen expression and the presentation of novel or previously unrecognised epitopes to the immune system leads to a loss of tolerance and initiation of disease. Subsequent generation of autoantibodies and generation of T cell responses may promote or propagate disease mechanisms.

## **1.9 Aims of study**

As outlined in this chapter, in view of the clinical heterogeneity of these conditions diagnosing patients as purely DM or PM is perhaps too broad. Many investigators have proposed new sets of criteria based on updated insights. Studies suggest that clinical stratification is more appropriate according to the MSA / MAA, as the detected autoantibody and clinical phenotype are closely associated (6, 139, 140).

The hypothesis for this body of research is that adult and juvenile IIM is associated with autoantibodies that recognise specific autoantigen systems, which define patients into homogeneous clinical phenotypes. To address this hypothesis the overall aim of this thesis is to describe a comprehensive clinical and serological study of adult and juvenile IIM, including the identification of novel autoantibody subsets.

### **Aim 1:**

To establish the prevalence and clinical features, in particular subtypes of recognised MSAs in the Royal National Hospital for Rheumatic Diseases (RNHRD) adult IIM cohort in comparison to previous published series.

### **Aim 2:**

To investigate and characterise novel autoantibody specificities and corresponding autoantigen targets in patients recruited to the RNHRD and UK Adult Onset Myositis Immunogenetic Collaboration (AOMIC) IIM cohorts.

### **Aim 3:**

To perform a clinical and serological study of children with JDM and JDM-overlap recruited to the UK and Ireland JDM Registry (UK JDM Cohort Study). To identify novel autoantibody-autoantigen systems, define clinical manifestations and thus markers of disease severity.

## **CHAPTER TWO**

### **PATIENTS AND METHODS**

#### **2.1 Patients and sera**

Subjects for the studies described in this thesis were recruited from three sources:

- a) The Royal National Hospital for Rheumatic Diseases Adult Connective Tissue Disease Clinic (RNHRD)
- b) The UK Adult Onset Myositis Immunogenetic Collaboration
- c) The Juvenile Dermatomyositis Registry and Repository, UK and Ireland

##### **2.1.1 Clinical diagnosis**

Adult IIM patients, aged 18 years of age or older at disease onset, were recruited through the RNHRD and the UK Adult Onset Myositis Immunogenetic Collaboration, (AOMIC) (9). JDM patients were recruited to the UK and Ireland JDM National Registry and Repository (JDRR) (24). The JDRR has recruited patients with juvenile-onset myositis, below the age of 16 years at disease onset and diagnosis. All adult PM/DM and JDM cases had probable or definite disease according to *Bohan and Peter* criteria (1, 2). For adult myositis/CTD-overlap cases recruited to AOMIC, patients were included if they fulfilled all of the following: a) met published criteria for their primary CTD or mixed connective tissue disease (MCTD) (11, 12, 141, 142); b) possessed at least two of four *Bohan and Peter* criteria; (c) possessed at least one MSA/MAA. JDM-scleroderma overlap (JDM-SSc) was defined as JDM children with a history of Raynaud's phenomenon, sclerodacty and other sclerodermatous skin changes (two or more of the above features).

Demographic and clinical data were recorded at diagnosis using standardised proformas (see Section 2.1.2 and 2.1.3). Serial clinical data was also available on patients attending the RNHRD and recruited through the JDRR. Data were stored using anonymous codes onto a central database. Serum and DNA samples were taken at the time of diagnosis and stored at -20°C until required.

##### **2.1.2 Adult-onset inflammatory myopathy clinical proformas**

See Chapter 2.5, Appendix I-III, pages 50-52.

##### **2.1.3 Juvenile-onset inflammatory myopathy clinical proformas**

See Chapter 2.5, Appendix IV, pages 53-60.

#### **2.1.4 Preparation and storage of sera**

Serum samples: approximately 5 ml of blood was venesected into a standard clotted sample tube. Once the blood had clotted, tubes were centrifuged at 3000 rpm for 10 min (Hereaus Labofuge 6000 Centrifuge). The serum fraction was extracted with a pipette into 500µl aliquots (adult samples) or 150µl aliquots (juvenile samples). Serum samples were stored at -20°C.

#### **2.1.5 Ethical approval**

Collection of data and blood from patients and controls was undertaken under the regulation of the local research ethics committees and informed consent was obtained according to the Declaration of Helsinki.

### **2.2 Tissue culture techniques**

All techniques were performed in class II tissue culture cabinets with sterile equipment. All reagents and media were purchased from Sigma, UK unless otherwise stated.

#### **2.2.1 Preparation of tissue culture media**

##### ***RPMI media***

One litre of RPMI-1640 medium (with NaHCO<sub>3</sub>, without L-glutamine) was supplemented with 10 ml 200 mM L-glutamine solution. Additionally, 100 ml foetal calf serum (heat inactivated) was also added for RPMI + FCS (10% foetal calf serum).

##### ***Preparation of dialysed foetal calf serum (FCS)***

Two sections of dialysis tubing were cut (1 metre each, medium width), placed in a beaker of tap water and left under a running tap for 3-4 hr. The tubing was transferred to 300 ml 3% (v/w) sodium sulphide at 80°C for 1 min, 300 ml distilled water at 60°C for 2 min and 300 ml 0.2% (v/v) sulphuric acid at room temperature for 1 min before being rinsed in distilled water. Tubing was finally placed in 200 ml 0.02% (w/v) sodium azide and stored at 4°C.

500 ml FCS was thawed at 37°C overnight and PBS (2 x 3 l) was cooled at 4°C overnight. The following day, lengths of treated dialysis tubing were rinsed in distilled water. FCS was transferred to the treated dialysis tubing, placed in 3 l cold PBS (prepared according to the manufacturer's instructions) and stirred at 4°C for 6 hr. The PBS was replaced with a further 3 l cooled PBS and stirred at 4°C for a

further 22 hr. Dialysed FCS was transferred to a tissue culture flask and sterile filtered through a 0.22µm filter unit. Filtered FCS was aliquoted into 50 ml tubes and stored at -20°C until required. The sterility of the dialysed FCS was tested by adding 2 ml to 20ml RPMI + FCS and incubating at 37°C for 48 hr.

#### ***Methionine free RPMI media (with dialysed foetal calf serum)***

One hundred ml RPMI-1640 medium (modified) (with NaHCO<sub>3</sub>, without methionine, cystine and L-glutamine) was supplemented with 62 µl 10% (w/v) cystine (prepared in 2M HCl and sterile filtered) and 1 ml 200mM L-Glutamine. Five ml dialysed FCS was added for methionine deficient RPMI + 5% dialysed FCS.

#### **2.2.2 Cell culture**

The non-adherent K562 cell line (human chronic myelogenous leukaemic cells) was obtained from European Collection of Cell Cultures (Salisbury, UK) (89121407). Cells were stored under liquid nitrogen at the University of Bath and transported to the cell culture facilities on dry ice.

#### ***Thawing K562 cell lines***

Vials of K562 cells in 90% FCS and 10% DMSO (dimethyl sulfoxide) were transferred to dry ice and transported to the BIRD laboratory, RNHRD. Aliquots were thawed at 37°C and added to 50 ml conical tubes containing 45 ml pre-heated RPMI + 10% FCS. Cells were incubated at 37°C for 10 min, before being centrifuged at 2000 rpm for 5 min (Heraeus Labofuge 6000). Supernatants were discarded and the pellets were resuspended in 25 ml pre-heated RPMI + 10% FCS. Cells were transferred to 60 ml flasks containing pre-heated RPMI + 10% FCS.

K562 cells were cultured and maintained in approximately 50 ml RPMI + 10% FCS, in vented, canted, tissue culture flasks. Cells were incubated in a 5% carbon dioxide humidifier incubator set at 37 °C.

### **2.3 Serological Methods**

Anti-nuclear autoantibody screening was performed by indirect immunofluorescence of NOVALite HEp-2 cells (NOVA Diagnostics, UK). Autoantibody specificity was further characterised by immunoprecipitation of radiolabelled cell extracts. All reagents and media were purchased from Sigma, UK unless otherwise stated.



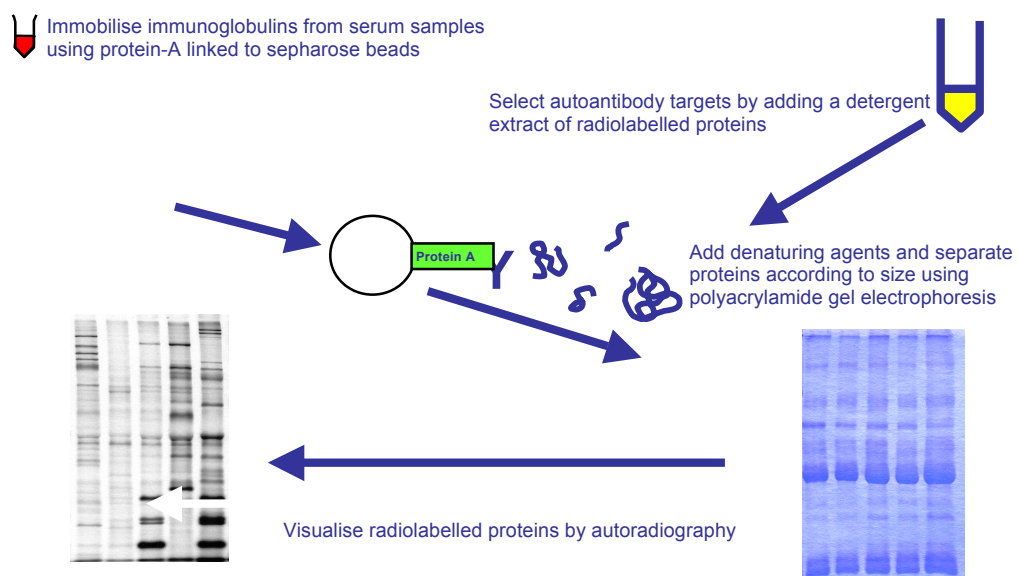
### 2.3.1 Indirect immunofluorescence

Serum was incubated against a fixed substrate (HEp-2 cells) to allow binding of autoantibodies to their respective antigen(s). A secondary goat polyclonal antibody conjugated to fluorescein was incubated against the substrate-antibody used at 1/150 dilution in the assay). Slides (binding site slides) were mounted and studied under a fluorescence microscope. The presence of ANA was recognised by particular patterns of fluorescence.

### 2.3.2 Radio immunoprecipitation (IPP) of protein targets

Serum was incubated with protein-A coated Sepharose beads to enable IgG present in the serum to bind to the protein-A via the Fc fragment. Beads were washed and supernatant removed (to remove any unbound material) and the beads incubated with radiolabelled K562 cell extract. Beads were washed further and the supernatant removed. Samples were re-suspended in Laemmli sample buffer. Finally, samples were boiled to remove and denature bound autoantigens from the Sepharose protein-A and radiolabelled proteins were separated by SDS-PAGE electrophoresis. Gels were analysed following autoradiography, protein targets were recognised by the characteristic gel fingerprint and the molecular weight of the band.

**Figure 3: Protein immunoprecipitation (IPP)**



#### ***Preparation of the of K562 cell extract with Methionine***

K562 culture supernatants from 6 x 150 ml flasks were transferred by pipette to 8 x 50 ml conical tubes. These were centrifuged at 2000 rpm for 5 min (Heraeus Labofuge 6000). Supernatants were sterilely removed and the pellets were pooled and re-suspended in 2 conical tubes containing 40 ml methionine free medium (without dFCS). Cells were centrifuged at 2000 rpm for 5 min and the supernatants sterilely removed. Pellets were re-suspended in 2 conical tubes each containing 40 ml methionine free medium (with FCS). A cell count (1/10 dilution) was completed on each tube.  $120 \times 10^6$  cells were added to 4 x 120 ml flasks and the flasks were topped up to 120 ml with methionine free medium (with dFCS). Cells were incubated at 37°C for 40 min. Two hundred and fifty µl radioactive methionine (Perkin-Elmer, UK EasyTag L-<sup>35</sup>S-Methionine, 5mCi (185 MBq) Stabilised Aqueous Solution) was added to each flask. The flasks were gently mixed and left at 37°C overnight. Culture supernatants were poured into 8 x conical tubes and were centrifuged at 2000 rpm for 5 min. The supernatants were discarded and the pellets pooled in 1 x conical tube containing 40 ml ice-cold Tris Buffered Saline (TBS) (10 mM Tris-HCl pH 7.4, 150 mM NaCl). A cell count was completed prior to centrifugation at 2000 rpm for 5 min. Supernatant was discarded and the cells were re-suspended in ice-cold immunoprecipitation (IPP) buffer (10 mM Tris-HCl pH 8.0, 500 mM NaCl 0.1% v/v Igepal) at a concentration of  $12 \times 10^6$  cells / ml. The extract was sonicated using a Soniprep 150 MSE sonicator over iced-water in 7.5 ml batches (amplitude 4.5, 1 min on, 20 sec off x 4 or until no viscous lumps remained). Sonicated extract was pooled into a fresh 50 ml conical tube and then aliquoted into 1.5 ml tubes. Extract was centrifuged at 120000 rpm for 20 min at 1°C (MSE Hawk 15/05). Supernatants were pooled, transferred to fresh tubes 1.5 ml tubes (1 ml aliquots); snap frozen in dry-ice and ethanol and stored at -80°C. Any radioactive contamination was cleaned up using lipsol and appropriate standard operating procedures.

#### ***Preparation of protein-A Sepharose beads***

Protein-A Sepharose beads were washed and hydrated in 40 ml IPP buffer by rotation at room temperature for 30 min. Hydrated Sepharose beads were centrifuged at 1100 rpm for 1 min (Heraeus Labofuge 6000). Supernatant was discarded and the pelleted beads were re-washed in 40 ml IPP buffer. Samples were re-centrifuged at 1100 rpm for 1 min, the supernatant was discarded and beads were re-suspended in IPP buffer at a concentration of 2 mg / ml.

### ***Immobilisation of Antibodies***

For each sample, 10 µl patient sera was mixed with 2 mg hydrated, washed protein-A Sepharose beads in 500 µl IPP buffer at room temperature for 30 min with end-over-end rotation. Autoantibody-coated Sepharose beads were pelleted at 2000 rpm for 1 min (MSE Hawk 15/05). Supernatant was discarded and the beads re-suspended in 1 ml ice-cold IPP buffer. The beads were washed a further 3 times by pelleting at 2000 rpm for 1 min, removing the supernatant and re-suspending in 1 ml ice-cold IPP buffer. After pelleting a final time at 2000 rpm for 1 min, the supernatant was discarded and the beads were resuspended in 380 µl ice-cold IPP buffer

### ***Immunoprecipitation of Antigen***

Radiolabelled K562 cell extract was thawed and 120 µl was added to each sample. Samples were mixed with end-over-end rotation at 4°C for 2 hr. Samples were centrifuged at 2000 rpm for 1 min and the supernatant discarded. Beads were washed 3 times in IPP (as outlined in the paragraph above) and once in 1 ml ice cold TBS. Beads were pelleted a final time at 2000 rpm for 1 min, the supernatant was discarded and beads were re-suspended in 50 µl SDS Laemmli sample buffer. Samples were vortexed briefly and stored at -80°C until required.

## **2.3.3 SDS polyacrylamide gel electrophoresis**

### ***Assembly of apparatus***

The Maxi gel electrophoresis equipment (Hoefer, UK) was assembled according to manufacturer's instructions. Glass plates were washed with water, dried and cleaned with methanol. One ml thick spacers separated Glass plates and 15 well combs were used.

### ***Preparation of gels***

A 10% resolving gel was used as the standard for this procedure (see Chapter 2, Appendix V for gel recipes). Resolving gel mixture was poured into the assembled gel apparatus up to 2 cm from the top of the glass plates. Water saturated butan-2-ol (approximately 300 µl) was layered onto the top of each gel. Gels were left for at least 15 min to polymerise prior to being rinsed with distilled water. The components for a 5% stacking gel were mixed and layered on top of the resolving gel to the top of the glass plates. Teeth combs were inserted to approximately 1 cm from the top of the resolving gel. The stacking gel was left to polymerise for approximately 15 min prior to the combs being removed and the wells rinsed with distilled water.

### ***Sample preparation***

Immunoprecipitated samples were taken out of the -80°C freezer to thaw. Samples were placed into the heating block and boiled at 95°C for 4 min, removed and briefly vortexed. Samples were centrifuged at 2000 rpm for 1 min (MSE Hawk 15/05) to pellet the beads.

### ***Gel loading and electrophoresis***

Wells were loaded with 25 µl supernatant per sample. On each gel, 1 well was reserved for 15 µl pre-stained molecular weight marker (Biorad, UK). When not all the wells were used samples were spaced evenly across the gel. Once loaded, the gel apparatus was fully assembled according to manufacture's instructions and transferred to the buffer chamber along with a cooling coil. Four L running buffer (25 mM Tris-base, pH7.4 190 mM Glycine and 0.1% w/v SDS) was added to the lower chamber and 500 ml running buffer was added to the upper chamber. A constant current of 60 mA was run until the dye from the sample buffer and markers had passed through the stacking gel layer (approximately 1 hr). The current was then increased to 120 mA run until the dye had passed through the resolving gel (approximately 3 hr).

### ***Gel fixing and drying***

Gels were removed from the glass plates and the stack layers discarded. The resolving gels were soaked in 0.5M sodium salicylate (enhancer) for 15 min in shallow trays. Salicylate solution was discarded; gels were rinsed in distilled water and placed in gel fix solution (methanol:water:glacial acid (4.5 :4.5:1 ratio)) for 30 min on a rocking platform. The fixative was drained off and the gels transferred onto blotting paper. The gels and blotting paper were placed on a gel dryer (Rapidry gel dryer ATTA Electrophoresis) and covered with cling-film. Gels were dried at 70°C for 70 min.

### ***Autoradiography***

Labelled proteins were analysed by autoradiography. In a dark room, dried gels were placed into X-ray cassettes and overlaid with Kodax Biomax film (mat side down). Cassettes were stored in a -80°C freezer for 1-4 weeks (time stored was dependant on the age of the radioactive preparation and the immunoreactivity of the autoantigen target). Development of films was undertaken manually in the dark room. Films were placed in 47 ml Kodax developer solution mixed with 460 ml distilled water for 2 min (until bands were exposed). Films were rinsed in distilled

water and placed into 47 ml Kodak fixer solution mixed with 460 ml distilled water for 5 min. Films were immersed in distilled water again and then left to dry at room temperature.

See Chapter 2, Appendix VI, Table 3: summary data of molecular weights / apparent molecular weight run on 10% SDS-PAGE of recognised myositis-specific autoantigen targets.

#### **2.3.4 Immunodepletion**

See 2.3.2 for basic IPP methods and reagents. This technique was used to establish whether autoantibodies that recognised a novel autoantigen of the same molecular weight (in specific patient cohorts) were targeting the same protein specificity. In summary, radiolabelled cell extracts were immunoprecipitated with prototype sera (sera known to contain the novel autoantibody specificity): Serum was incubated with protein-A bound Sepharose beads and incubated with radiolabelled K562 cell extract. On this occasion, the pelleted beads bound to the autoantibody-autoantigen complex were removed and the supernatant (unbound cell extract) preserved. This procedure was repeated to ensure complete depletion of the target antigen from the cell extracts. Reference depleted cell extracts were then used for further immunoprecipitations with other autoantibody positive sera.

In detail, for each sample, duplicate tubes each containing 10 mg hydrated protein-A Sepharose beads (prepared as Section 2.3.2) and 50 µl reference sera in 1 ml IPP buffer were mixed with end-over-end rotation at room temperature for 30 min. Autoantibody-coated Sepharose beads were washed 4 x in IPP buffer (see section 2.3.2). After the beads were pelleted for the final time, supernatants were discarded. One tube for each sample (A) was placed on ice, whilst 150 µl [<sup>35</sup>S]-methionine-labelled K562 cell extract and 380 µl IPP buffer was added to the remaining tube (B). Tube B was mixed with end-over-end rotation at 4°C for 2 hr after which the supernatant was transferred to tube A. This was mixed at 4°C for a further 2 hr. The supernatant from tube A was transferred to a fresh tube (C) and stored at -80°C. Immunoprecipitations using positive sera and either 120µl control [<sup>35</sup>S]-methionine-labelled cell extract or the immunodepleted supernatants (C) were completed and subject to SDS-PAGE electrophoresis as described in Section 2.3.2.

### **2.3.5 Immunoprecipitation Western Blotting**

Immunoprecipitation Western Blotting (IPP-WB) was used to determine the identify of specific autoantigens recognised by autoantibodies. Autoantigens were selected from patient's sera by immunoprecipitation and SDS-PAGE. Proteins were transferred to nitrocellulose membrane by immunoblotting. Membranes were incubated in a protein rich solution to block any remaining non-specific binding sites and probed with commercial antibody to a specific protein. Membranes were incubated with an appropriate secondary labelled anti-immunoglobulin followed by a conjugated staining substrate.

#### ***Preparation of unlabelled K562 cell extract***

K562 cultures were grown to confluence in 6 flasks each containing 150 ml complete RPMI with FCS (see section 2.2.1). Supernatants were transferred to 8 x 50 ml conical tubes and centrifuged at 2000 rpm for 5 min (Heraeus Labofuge 6000). Supernatants were removed and the pellets resuspended in 2 x conical tubes each containing 40 ml ice-cold TBS. A cell count (1/10 dilution) was completed on each tube. Cells were centrifuged at 2000 rpm for 5 min, the supernatants were discarded and the pellets combined in a single conical tube containing ice-cold IPP buffer at a concentration of  $15 \times 10^6$  cells / ml. Extract was sonicated using a Soniprep 150 MSE sonicator over iced-water in 7.5 ml batches (amplitude 4.5, 1 min on, 20 sec off x 4 or until no viscous lumps remained). Sonicated extract was pooled into a fresh 50 ml conical tube and aliquoted into 1.5 ml tubes. Extract was centrifuged at 120000 rpm for 20 min at 1°C (MSE Hawk 15/05). Supernatants were pooled, transferred to fresh 1.5 ml tubes (1.2 ml aliquots); snap frozen in dry-ice and ethanol and stored at -80°C.

#### ***Antigen preparation for IPP WB***

See 2.3.2 for IPP methods and reagents. Samples, each containing 40 µl serum and 2 mg hydrated protein-A Sepharose beads (prepared as Section 2.3.2) in 500 ml IPP buffer were mixed with end-over-end rotation at room temperature for 30 min. Autoantibody-coated Sepharose beads were centrifuged at 2000 rpm for 1 min (MSE Hawk 15/05), supernatant was discarded and beads re-suspended in 1 ml 0.2M tri-ethanolamine pH 8.1. Beads were washed a second time (centrifuged at 2000 for 1 min and re-suspended in tri-ethanolamine) prior to being centrifuged again and re-suspended in 5 mM bis-(sulphosuccinimidyl)-suberate cross-linker (Perbio, UK) in 1 ml 0.2M tri-ethanolamine. Beads were mixed with end-over-end rotation at room temperature for 30 min and centrifuged at 2000 rpm for 1 min.

Beads were re-suspended in 1 ml 50 mM Tris-Cl pH 7.5 and mixed with end-over-end rotation at room temperature for 15 min. Beads were centrifuged at 2000 rpm for 1 min and washed three times in PBS and two times in ice-cold IPP buffer. Beads were incubated in 1 ml unlabelled K562 cell extract with end-over-end rotation for 1 hr at 4°C, centrifuged at 2000 rpm for 1 min and resuspended in 1 ml fresh K562 unlabelled cell extract. Samples were incubated for a further 1 hr at 4°C with end-over-end rotation. Beads were washed 4 times in ice-cold IPP buffer and once in TBS. Samples were finally re-suspended in 80 µl SDS sample buffer and stored at -80°C until required.

### ***SDS-PAGE and transfer***

Proteins were fractionated using by 10% SDS-PAGE as per standard methods (see 2.3.3), however samples were loaded at 80 µl per lane. During the electrophoresis, blotting sheets and nitrocellulose membranes were soaked in Transfer Buffer pH7.4 (40mM tris, 20 mM Sodium acetate, 2mM EDTA, 20% v/v methanol and 0.1% w/v SDS). Once electrophoresis had been completed, gels were placed in the immunoblotting transfer apparatus according to manufacturers' instructions. 'Transfer sandwiches' were prepared as scotchbrite, blotting paper, nitrocellulose membrane, gel, blotting paper and scotchbrite. The surfaces of the 'transfer sandwiches' were smoothed over to remove any air-bubbles. The 'transfer sandwiches' were placed into the tank apparatus that was filled with Transfer Buffer. A cooler system was inserted and proteins were transferred from the gels to the nitrocellulose membranes using a constant current at 220 mA for 4.5 hr. Nitrocellulose sheets were removed and the corners were marked. Membranes were rinsed in distilled water and were Ponceau stained for 5 min. Membranes were rinsed again for 4 x 15 sec in distilled water and air-dried overnight. Membranes were stored in a sealed container at 4°C until required.

### ***Immunoblotting and staining***

Membranes were washed in PBS-0.05%Tween for 5 min and blocked for 90 min in block solution (5% milk powder in PBS-0.05%Tween, pH7.4). Membranes were incubated with commercial primary antibody (diluted in block solution accordingly to manufacturer's instructions) for 90 min. Membranes were washed 5 x 5 min in PBS-0.05%Tween and re-incubated with the corresponding secondary antibody (commercial anti-IgG) at a 1:20000 dilution for 90 min. Membranes were washed 5 x 5 min in PBS-0.05%Tween and finally incubated in substrate stock for 10-20 min. Membranes were rinsed with distilled water for 5 min and air-dried overnight.

## **2.4 Statistical analysis**

Clinical and autoantibody associations were derived from 2×2 contingency tables using the chi-squared test with Yates' continuity correction, or two-tailed Fisher's exact test where individual cells valued five or less. Where significant, data were expressed as odds ratios (OR) with exact 95% confidence intervals (CI). Where data was not normally distributed the Mann-Whitney-U test was used to compare continuous data. Median values (inter-quartile ranges) were expressed where appropriate.  $P$  values ( $P_{\text{corr}}$ ) were adjusted using the Bonferroni correction when comparing clinical associations. Uncorrected probabilities are presented for possible clinico-serological associations.  $P$  values <0.05 were considered significant. *SPSS for Windows (version 14)* was used to perform statistical analysis.



## **2.5 Appendix**

- I. RNHRD Clinical proforma (Page 50)
- II. AOMIC Clinical Proforma (Page 51)
- III. Autocure Clinical Proforma (Page 52)
- IV. JDRR clinical Proforma (forms 1 and 2) (Pages 53-60)
- V. Reagents and gel recipes (Pages 61-63)
- VI. Table 3: Summary data of molecular weights / apparent molecular weight run on 10% SDS-PAGE of recognised myositis-specific autoantigen targets (Page 64)

**Permission to reproduce AOMIC, Autocure and JDRR proformas granted by:**

\*Dr. Hector Chinoy, Consultant Senior Lecturer, University of Manchester

\*\*Prof. Lucy Wedderburn, Scientific Director, JDRR, University College London



## MYOSITIS GENETIC STUDY CLINICAL & LABORATORY PROFORMA

### PATIENT DETAILS (complete or ring)

NAME: .....

Sex: M / F Hospital No.....

Ethnic origin: Caucasian/ Negroid / Asian / Oriental  
Mixed (specify.....)  
Other (specify.....)

Current age:.....

Age at myositis onset:.....

### MYOSITIS DIAGNOSTIC GROUP (tick or ring)

Polymyositis (PM) [ ]  
Dermatomyositis (DM) [ ]  
PM as part of other CTD [ ]

(Specify: SLE / SSs / MCTD / Overlap / Undifferentiated)

**SYMPTOM / LESION INDEX** - We will only include definite PM/DM, according to the diagnostic criteria highlighted, so please tick which abnormalities ever present:

1. Skin Lesions
  - Heliotrope rash [ ]
  - Gottron's sign [ ]
  - Violaceous rash on elbows/knees [ ]
2. Systemic inflammatory signs [ ]
3. Non-destructive arthropathy [ ]
4. Muscle-specific antibody [ ]
5. Elevated skeletal muscle enzymes [ ]
6. Myalgia at rest or on contraction [ ]
7. Proximal muscle weakness [ ]
8. Myopathic EMG [ ]
9. Myositis on biopsy [ ]

If patient has lung fibrosis as part of their PM, tick below.  
Lung fibrosis (clinically and/or on PFTs) [ ]

### DIAGNOSTIC CRITERIA FOR PM/DM

- 1) Skin lesions:
  - Heliotrope rash (violaceous erythema on upper eyelids),
  - Gottron's sign (violaceous keratotic papules on extensor aspects of fingers,
  - Violaceous raised rash over elbows or knees.
- 2) Systemic inflammatory signs (elevated ESR/CRP, pyrexias, weight loss etc)
- 3) Non-destructive arthritis
- 4) Muscle specific antibodies, e.g Anti Jo-1 etc
- 5) Elevated skeletal muscle enzymes
- 6) Myalgia at rest or with contraction
- 7) Proximal muscle weakness
- 8) Myopathic EMG
- 9) Myositis on muscle biopsy.

#### For diagnosis of definite PM:

At least four from item list 2 – 9.

#### For diagnosis of definite DM:

At least one feature from item 1), plus at least four from item list 2 – 9.

(From Tanimoto et al. Classification criteria for PM/DM. J. Rheumatol 1995, 22: 668-674)

**UNDERLYING MALIGNANCY** - If your patient has PM/DM, in association with a proven malignancy, could you please specify site & tissue type if known

.....  
.....

Patient's Consultant:.....

Hospital:.....

Date form completed:.....

Signature:.....

RGC/2000

AUTOCURE MYOSITIS CLINICAL & LABORATORY PROFORMA	
<b>1. CENTRE &amp; PATIENT DETAILS</b> (complete / attach sticker):	
HOSPITAL (Name/Code):.....	
PATIENT (Name):.....	
Gender (ring): M / F	
NHS/Hospital N <sup>o</sup> :.....	
Ethnicity: Caucasian/ Afro-Carrib/ African/ Asian/ Oriental Mixed (specify):..... Other (specify):.....	
Date of birth:.....	
Month/Year of myositis onset:.....	
Month/Year of myositis diagnosis:.....	
Ht (cm)..... Wt (kg)..... Ever smoked?: YES / NO	
<b>2. MYOSITIS DIAGNOSTIC GROUP</b> (please tick one of a-d):	
a) Polymyositis (PM)	<input type="checkbox"/>
b) Dermatomyositis (DM)	<input type="checkbox"/>
c) Amyopathic Dermatomyositis (ADM)	<input type="checkbox"/>
d) Inclusion Body Myositis (please complete IBM-specific form) (IBM)	<input type="checkbox"/>
ADDITIONALLY, if overlap with CTD: SLE/ Sjögren's/ MCTD/ SSc/ RA (ring) (Overlap) <input type="checkbox"/>	
other (specify):.....	
<b>3. CASE VERIFICATION</b> (please tick all that apply):	
1. Proximal muscle weakness	<input type="checkbox"/>
2. Myositis on biopsy	<input type="checkbox"/>
3. Elevated skeletal muscle enzymes	<input type="checkbox"/>
4. Myopathic EMG	<input type="checkbox"/>
5. Characteristic skin lesions	<input type="checkbox"/>
<b>4. OTHER CLINICAL SIGNS</b> (please tick):	
- Heliotrope rash	<input type="checkbox"/>
- Gottron's papules	<input type="checkbox"/>
- Violaceous rash on elbows/knees	<input type="checkbox"/>
- V sign	<input type="checkbox"/>
- Shawl sign	<input type="checkbox"/>
- Periungual erythema	<input type="checkbox"/>
- Mechanics' hands	<input type="checkbox"/>
- Raynaud's	<input type="checkbox"/>
- Arthritis	<input type="checkbox"/>
- Dysphagia	<input type="checkbox"/>
- Arrhythmias	<input type="checkbox"/>
JV/RGC/HC/2009	
<b>5. AUTO-ANTIBODY DETECTION</b> (please ring):	
Myositis specific/associated autoantibody	YES / NO
Specify:.....	
Non myositis-specific antibodies	YES / NO
Specify:.....	
<b>6. INTERSTITIAL LUNG DISEASE</b> (please ring)	
(attributable to myositis)	YES / NO
Proven by (ring):	PFTs / CXR / HRCT
Month/year of diagnosis: .....	
<b>7. MALIGNANCY:</b> (regardless of whether you consider the myositis to be cancer-associated)	
Month/year of cancer diagnosis:.....	
Site/tissue type:.....	
<b>8. ENVIRONMENTAL FACTORS:</b> (please ring/complete)	
?Statin / fibrate use prior to onset of myositis	YES / NO
(specify drug if known):.....	
Patient's view of what triggered myositis (specify):.....	
Exposures: Asbestos/Silica/Fibreglass/Solvents/Coaldust other (specify):.....	
Patient's Physician (print):.....	
Date form completed (print):.....	
<b>Bohan and Peter Diagnostic Criteria for Adult PM/DM<sup>1</sup></b>	
1) Symmetrical weakness of limb-girdle muscles and/or anterior neck flexors.	
2) Muscle biopsy evidence typical of myositis.	
3) Elevation of serum skeletal muscle enzymes.	
4) Typical EMG features of myositis.	
5) Typical DM rash, including: - Heliotrope rash: violaceous erythema on upper eyelids. - Gottron's papules: violaceous keratotic papules on extensor aspects of fingers. - Violaceous rash over extensor aspects elbows/knees.	
Exclusion criteria: Absence of other forms of myopathy, e.g. inclusion body, metabolic, inherited or infectious forms.	
Probable / definite PM: at least 3 of items 1-4 +ive. Probable / definite DM: item 5 & at least 2 of items 1-4 +ive.	
Adult PM/DM	Must fulfill probable or definite Bohan & Peter criteria (see highlighted box).
Adult PM/DM-Overlap	Must fulfill criteria for their main CTD <sup>2-6</sup> and at least 2 of items 1-4 and have at least one myositis specific / associated autoantibody.
JDM	Attending physician's diagnosis accepted (tick which criteria were satisfied for diagnosis).
Amyopathic DM	Must satisfy Sontheimer's criteria <sup>7</sup> .
IBM	Must satisfy Griggs's criteria <sup>8</sup> .
<b>References:</b>	
1. PM/DM, Bohan A, Peter JB. N Engl J Med 1975;292: 344-7; 2. SLE, Tan EM et al. Arthritis Rheum 1982;25:1271-7; Hochberg MC. Arthritis Rheum 1997;40:1725; 3. SSc, Diagnostic and Therapeutic Criteria Committee. Arthritis Rheum 1980;3:581-90; 4. RA, Arnett FC et al. Arthritis Rheum 1988;31:315-4; 5. Sjögren's, Vitali C et al. Arthritis Rheum 1993;36:340-7; 6. MCTD, Alarcon-Segovia D. Clin Dermatol 1994;12:309-18; 7. Amyopathic, Sontheimer RD. J Am Acad Dermatol 2002; 46:626-36; 8. IBM, Griggs RC et al. Ann Neurol 1995;38:705-13.	

**JUVENILE DERMATOMYOSITIS COHORT BIOMARKER STUDY AND REPOSITORY  
FORM 1 – INITIAL PRESENTATION INFORMATION**

**A. DEMOGRAPHICS:**

DATE OF VISIT:

PATIENT REF NO:

Patient status\*:

Male/Female

DOB:

Ethnicity: 01 02 03 04 05 06 07 08 09

(Please see guidelines for codes. If 04 or 09, please specify)

Postcode:

Occupation of parents:



Please tick

☐

Has consent/assent form for the registry and repository been completed?

\*1=Initial visit/diagnosis at GOS/MDX

2=GOS/MDX primary institution for JDM care after previous diagnosis elsewhere

3=Seen for occasional care/shared care only (i.e. GOS/MDX not primary centre)

4=Other (please specify)

**B. HISTORY UP TO DIAGNOSIS:**

**1. Background data**

DIAGNOSIS: \_\_\_\_\_

**I. General symptoms:**

Absent Present Not known

Rash 0 1 9  
Weakness 0 1 9  
Fever 0 1 9  
Alopecia 0 1 9  
Weight loss 0 1 9  
Fatigue 0 1 9  
Mouth ulcers 0 1 9  
Headache 0 1 9  
Irritability 0 1 9  
Raynauds 0 1 9

y):  
yyyy

**II. Musculoskeletal:**

Absent Present Not known

Myalgia 0 1 9  
Joint pain 0 1 9  
Jt stiffness 0 1 9  
Jt swelling 0 1 9  
Dyspnoea 0 1 9  
Dysphonia 0 1 9  
Dysphagia 0 1 9

At diagnosis if available

Height: \_\_\_\_\_cm

Weight: \_\_\_\_\_kg

BP: \_\_\_\_\_/\_\_\_\_\_

Not known  
9

Diarrhoea 0 1 9  
Melaena 0 1 9  
Haematuria 0 1 9  
Facial/body swelling 0 1 9

<b>3. Past medical history</b>	<i>Absent</i>	<i>Present</i>	<i>Please specify where details known/relevant</i>
Specific viral infection	0	1	
Other autoimmune disease	0	1	
Other significant diagnosis	0	1	
Trauma/injury	0	1	
Immunisation in last 6/12	0	1	
School absence	0	1	
Stopped PE/Sport	0	1	
Respiratory problems	0	1	
GI problems	0	1	
Neurological involvement	0	1	
Medication related problems	0	1	
<b>FH of autoimmune disease</b>	0	1	

**C. EXAMINATION FINDINGS AT DIAGNOSIS:**

<u>I. Skin:</u>	<i>Absent</i>	<i>Present</i>	<i>Not known</i>
Gottroons papules	0	1	9
Ulceration	0	1	9
Oedema	0	1	9
Calcinosis	0	1	9
Nailfold changes	0	1	9
Lipoatrophy	0	1	9
Other:			
Specify/describe: _____			

	<i>Absent</i>	<i>Present</i>
<u>II. Arthritis</u>	0	1
<u>III. Muscle weakness</u>	0	1
<u>IV. Other:</u>	0	1
Specify _____		

**D. CLINICAL COURSE SINCE DIAGNOSIS: ( Tick this box and go to form 2 if patient recruited at diagnosis )**

<u>I. General symptoms:</u>			
	<i>Absent</i>	<i>Present</i>	<i>Not known</i>
Rash	0	1	9
Weakness	0	1	9
Fever	0	1	9
Alopecia	0	1	9
Weight loss	0	1	9
Fatigue	0	1	9
Mouth ulcers	0	1	9
Headache	0	1	9
Irritability	0	1	9
Raynauds	0	1	9

<u>II. Musculoskeletal:</u>			
	<i>Absent</i>	<i>Present</i>	<i>Not known</i>
Myalgia	0	1	9
Joint pain	0	1	9
Jt stiffness	0	1	9
Jt swelling	0	1	9
Dyspnoea	0	1	9
Dysphonia	0	1	9
Dysphagia	0	1	9

<u>III. Systemic features:</u>			
	<i>Absent</i>	<i>Present</i>	<i>Not known</i>
Chest pain	0	1	9
Abdo pain	0	1	9
Diarrhoea	0	1	9
Melaena	0	1	9
Haematuria	0	1	9
Facial/body swelling	0	1	9
Calcinosis	0	1	9

### E. INVESTIGATION RESULTS AT DIAGNOSIS:

0=normal, 1=abnormal, 9=not done: Please tick the appropriate box for each test result.

INVESTIGATION	DATE	RESULT			Value	FURTHER INFORMATION Please record any significant information
		0	1	9		
<b>Haematology</b>						
Hb						
WBC						
Platelets						
WBC Diff: Neut						
WBC Diff: Lymph						
ESR						
<b>Biochemistry</b>						
CRP						
Urea						
Creatinine						
CK						
LDH						
ALT						
AST						
Albumin						
Other (specify):						
<b>Immunology</b>						
RF						
ANA						
ENA						
dsDNA						
Other antibodies (specify):						
C3						
C4						
IgG						
IgA						
IgM						
ASOT						
<b>Radiology</b>						
MRI						
Muscle Ultrasound						
DEXA: T Score						
DEXA: Z Score						
<b>Lung Function</b>						
VC						
FEV1						
DLCO						
<b>Other</b>						
EMG						
Muscle Biopsy						
Skin Biopsy						
Barium Swallow						
Echocardiogram						
Urine Dipstick						
Other (specify):						

## **F. THERAPY INFORMATION SINCE DIAGNOSIS**

### **I. CORTICOSTEROIDS**

	<i>No</i>	<i>Yes</i>	<i>Not known</i>
Oral steroids	0	1	9
IV Steroids	0	1	9

### **III. NSAIDs**

	<i>No</i>	<i>Yes</i>	<i>Not known</i>
Naproxen	0	1	9
Other (specify):	0	1	
_____			

### **II. DMARDs**

	<i>No</i>	<i>Yes</i>	<i>Not Known</i>
Methotrexate	0	1	9
Cyclosporin A	0	1	9
Azathioprine	0	1	9
Cyclophosphamide	0	1	9
Hydroxychloroquine	0	1	9
IV-Immunoglobulin	0	1	9
Etanercept	0	1	9
Infliximab	0	1	9
Other (specify):	0	1	
_____			

### **IV. OTHER**

	<i>No</i>	<i>Yes</i>	<i>Not known</i>
Calcium/Vit D	0	1	9
Folic acid	0	1	9
Plasmaphoresis	0	1	9
Other (specify):	0	1	
_____			

## **G. PHYSIOTHERAPY/OCCUPATIONAL THERAPY**

	<i>No</i>	<i>Yes</i>	<i>Not known</i>
Dry land	0	1	9
Hydrotherapy	0	1	9
Splinting	0	1	9
Other	0	1	Specify:
_____			

## **H. OTHER INFORMATION**

*No*      *Yes*  
0        1

Specify:

<b>Signature:</b>	<b>Position:</b>
<b>Print name:</b>	<b>Date:</b>



# JUVENILE DERMATOMYOSITIS COHORT BIOMARKER STUDY AND REPOSITORY

## FORM 2: CLINIC VISIT FORM

Patient Reference No:

Date of visit:

### **A. HISTORY SINCE LAST CLINIC VISIT** **(or in last 3 months – whichever is most recent)**

#### **1. Clinical History**

<u>I. General Symptoms</u>	<i>Absent</i>	<i>Present</i>
Rash	0	1
Weakness	0	1
Fever	0	1
Alopecia	0	1
Weight loss	0	1
Fatigue	0	1
Mouth ulcers	0	1
Headache	0	1
Irritability	0	1
Raynauds	0	1

<u>II. Musculoskeletal</u>	<i>Absent</i>	<i>Present</i>
Myalgia	0	1
Joint Pain	0	1
Joint stiffness	0	1
Joint Swelling	0	1
Dyspnoea	0	1
Dysphonia	0	1
Dysphagia	0	1

<u>III. Systemic Features</u>	<i>Absent</i>	<i>Present</i>
Chest pain	0	1
Abdo pain	0	1
Diarrhoea	0	1
Melaena	0	1
Haematuria	0	1
Facial/body swelling	0	1

#### **2. Measurements**

Current height: \_\_\_\_\_cm

Current weight: \_\_\_\_\_Kg

BP: \_\_\_\_\_

#### **3. Family History**

No Yes

Any new family history in the last 3 months: 0 1  
Specify:

### **B. EXAMINATION FINDINGS**

<u>I. Skin</u>	<i>Absent</i>	<i>Present</i>
Gottrons papules	0	1
Ulceration	0	1
Lipoatrophy	0	1
Oedema	0	1
Nailfold changes	0	1
Calcinosis	0	1
Other	0	1
Specify/describe: _____		

<u>II. Distribution of rash</u>	<i>Absent</i>	<i>Present</i>
Periorbital	0	1
Periungal	0	1
Trunk	0	1
Small joints	0	1
Large joints	0	1
Other	0	1
Specify: _____		

**III. Joints**

	<i>Absent</i>	<i>Present</i>	<i>Please specify which joints are affected</i>
Arthritis	0	1	_____
Pain on motion	0	1	_____
Joints with limited ROM	0	1	_____
Contractures	0	1	_____

**IV. Manual Muscle Testing**

Neck flexors	0	1	2	3	4	5	9
Shoulder abductors	0	1	2	3	4	5	9
Elbow flexors	0	1	2	3	4	5	9
Wrist Extensors	0	1	2	3	4	5	9
Hip Extensors	0	1	2	3	4	5	9
Hip Abductors	0	1	2	3	4	5	9
Knee Extensors	0	1	2	3	4	5	9
Ankle Dorsiflexors	0	1	2	3	4	5	9

\*0=no muscle action, 1=flicker of muscle action, 2=muscle action with gravity counterbalance, 3=muscle action against gravity, 4=muscle action against gravity with some resistance, 5=full muscle strength, (9=not done)

**V. Oedema**

	<i>Absent</i>	<i>Present</i>
Periorbital/facial	0	1
Limb	0	1
Trunk	0	1

**VI. Abdomen**

	<i>Absent</i>	<i>Present</i>
Abdominal masses	0	1
Tenderness	0	1
Hepatomegaly	0	1
Splenomegaly	0	1

**VII. Respiration**

*Please circle one:*

0      1      2      3      4      5

0=normal, 1=SOBE, 2=tachypnoea, 3=accessory muscle use, 4=requires oxygen, 5=ventilated

**VIII. Other**

	<i>Absent</i>	<i>Present</i>	<i>Please specify</i>
Calcinosis	0	1	_____
Eyes (glaucoma/cataracts)	0	1	_____
Other	0	1	_____

**IX. Physician Global Assessment**

*Disease least active*

0

*Disease most active*

10

**C. PHYSIOTHERAPY/OCCUPATIONAL THERAPY**

	<i>No</i>	<i>Yes</i>	
Dry Land	0	1	Daily/weekly/other
Hydrotherapy	0	1	Daily/weekly/other
CMAS done Today	0	1	Score: ____/53
CHAQ done today	0	1	Score: ____/3
CHQ done today	0	1	Physical score: ____ Psychological score: ____
Parents VAS	0	1	Score: ____
Splinting	0	1	Specify: _____
Other	0	1	Specify: _____

**D. INVESTIGATION RESULTS:** 0=normal, 1=abnormal, 9=not done - tick the appropriate box for each result

INVESTIGATION	DATE	RESULT			VALUE	FURTHER INFORMATION Please record any significant information
		0	1	9		
<b>Haematology</b>						
Hb						
WBC						
Platelets						
WBC Diff: Neut						
WBC Diff: Lymph						
ESR						
<b>Biochemistry</b>						
CRP						
Urea						
Creatinine						
CK						
LDH						
ALT						
AST						
Albumin						
Other (specify):						
<b>Immunology</b>						
RF						
ANA						
ENA						
dsDNA						
Other antibodies (specify):						
C3						
C4						
IgG						
IgA						
IgM						
ASOT						
<b>Radiology</b>						
MRI						
Muscle Ultrasound						
DEXA: T Score						
DEXA: Z Score						
<b>Lung Function</b>						
VC						
FEV1						
DLCO						
<b>Other</b>						
EMG						
Muscle Biopsy						
Skin Biopsy						
Barium Swallow						
Echocardiogram						
Urine Dipstick						
Other (specify):						

**E. THERAPY INFORMATION (since last visit)**

<b>I. Corticosteroids</b>					
	No	Yes	Current Dose	Reducing Dose?	
Oral Steroids	0	1	_____mg/day	Yes	No
IV Steroids	0	1			

<b>II. DMARDs</b>					
	No	Yes	Dose and type administered	Dates if started or stopped in the last 3 months	
Methotrexate	0	1	_____	_____	
Cyclosporin A	0	1	_____	_____	
Azathioprine	0	1	_____	_____	
Cyclophosphamide	0	1	_____	_____	
Hydroxychloroquine	0	1	_____	_____	
IV-IG	0	1	_____	_____	
Plasmapheresis	0	1	_____	_____	
Etanercept	0	1	_____	_____	
Infliximab	0	1	_____	_____	

<b>III. NSAIDs</b>				
	No	Yes	Dose on day attending clinic	
Ibuprofen	0	1	_____	
Naproxen	0	1	_____	
Other	0	1	_____	
Specify: _____				

<b>IV. Other Drugs</b>				
	No	Yes	Dose on day attending clinic	
Calcium/Vitamin D	0	1	_____	
Folic acid	0	1	_____	
Gastroprotectant	0	1	_____	
Vasodilator	0	1	_____	
Anti-HT	0	1	_____	
Other	0	1	_____	
Specify: _____				

**F. RESEARCH INVESTIGATION SAMPLES**

Were the following samples taken today: No Yes

Blood:

Lab white bottle (hep)	0	1
Clotted bottle	0	1

**G. OTHER INFORMATION**

Specify:

Signature:	Position:
Print Name:	Date:

## V. Reagents

- RPMI x1 liquid (Sigma)
- 10% FCS or FBS (Sigma)
- L-glutamine, 200mM (Sigma)
- Methionine free medium; solid (Sigma)
- L-leucine; solid (Sigma)
- L-lysine; solid (Sigma)
- Sodium bicarbonate (Tissue culture grade, Sigma)
- S<sup>35</sup> methionine (Amersham Biosciences (GE Healthcare) AG 1095 5mCi)
- Protein A or protein G sepharose beads (Sigma)
- Immunoprecipitation (IPP) buffer  
(10mM Tris/HCl, pH 8.0, 500mM NaCl, 0.1% v/v Igepal; previously known as Nonidet P-40)
- Tris buffered saline (TBS) buffer  
(10mM Tris/HCl, pH 7.4 + 150mM NaCl)
- Sodium Phosphate (crosslinking for MS)  
0.1M Sodium Phosphate pH8.1 (0.1M NaH<sub>2</sub>PO<sub>4</sub> to 500ml 0.1M Na<sub>2</sub>HPO<sub>4</sub> until pH=8.1)
- Tris-Cl (crosslinking stop buffer)  
1.0M Tris base pH7.5
- Triethanolamine (crosslinking reaction buffer)  
0.2M Triethanolamine pH8.1
- Laemmli sample Buffer (SDS-PAGE)  
Sigma S3401
- Western Blot Substrate  
Sigma B1911
- SDS-PAGE Marker  
Biorad Kaleidoscope Marker - 161-0324
- Blocking powder (Western Blot)  
5% Sainsburys semi-skimmed milk powder, pH 7.4 (with NaOH)
- PBS-Tween  
PBS with 0.05% Tween
- Phosphate Buffered Saline  
Sigma P4417 (1 tablet per 200 ml water)
- Western Blot Transfer Buffer  
40 mM Tris, 20 mM Sodium Acetate and 2 mM EDTA  
pH 7.4 with acetic acid  
Trizma base (Sigma)

## **V. Reagents**

- Glycine
- Sodium chloride (BDH)
- Conc. HCl (Sigma)
- Glacial acetic acid (Sigma)
- PBS tablets (Sigma)
- Acrylamide/bis-acrylamide, 30% solution (Sigma)
- Ammonium persulphate (Sigma)
- Kalidescope markers, broad range containing glycerol, protein and bromophenol blue (Bio-Rad)
- TEMED (Sigma)
- Sodium dodecyl sulphate (SDS), (Sigma)
- Sample buffer containing mercaptoethanol, glycerol, bromophenol blue, tris and SDS (Sigma)
- Methanol (Sigma)
- Butan-2-ol (Sigma)
- Sodium salicylate (Sigma)
- DMSO (Sigma)
- Trypan blue (Sigma)
- Conc. Sulphuric acid (BDH)
- Sodium sulphide (Sigma)
- Sodium azide (Sigma)
- pH buffer standards (Lab 3)

## V. Gel recipes

Percentage acrylamide in gel mixture (volumes are per gel)

Vol (ml)	5%	8%	9%	10%	12.5%	15%
Water	11.8	14.5	13.5	12.5	10	7.5
Lower gel buffer		7.5	7.5	7.5	7.5	7.5
Upper gel buffer	5					
Acrylamide mixture	3.2	8	9	10	12.5	15
AMPS	0.1	0.21	0.175	0.15	0.11	0.09
TEMED	0.03	0.06	0.04	0.03	0.02	0.02
Total volume	~20	~30	~30	~30	~30	~30

30% (w/v) acrylamide:bisacrylamide (30:5:1)

Lower gel buffer stock: 1.5M Tris/HCl, pH 8.8 + 0.4% (w/v) SDS

Upper gel buffer stock: (\*5% upper gels in this procedure) 0.5M Tris/HCl pH 6.8 + 0.4% (w/v) SDS

10% (w/v) ammonium persulphate

**VI. Table 3: Summary data of molecular weights / apparent molecular weight run on 10% SDS-PAGE of recognised myositis-specific autoantigen targets**

	<b>Autoantigen</b>	<b>Molecular Weight (kDa)</b>	<b>Runs at (10% SDS-PAGE) (kDa)</b>
Jo-1	Histidyl-tRNA synthetase	57.4	52
PL-7	Threonyl-tRNA synthetase	83.4	80
PL-12	Alanyl-tRNA synthetase	106.7	110
KS	Asparaginyl-tRNA synthetase	62.9	65
EJ	Glycyl-tRNA synthetase	83.1	75
OJ	Multi-enzyme complex		
	Leucyl-tRNA synthetase	125.0	130
	Isoleucyl-tRNA synthetase	145.0	150
	Glutaminyl-tRNA synthetase	160.0	170
Ha	Tyrosyl-tRNA synthetase	59	62
SRP	Signal recognition particle	262.3	240
Mi-2	SNF2 super-family helicase	55.7	54



## CHAPTER THREE

### RESULTS

#### RNHRD clinico-serological study of myositis-specific autoantibodies in adult-onset inflammatory myositis

##### 3.1 Introduction

There is increasing evidence that patients with adult-onset idiopathic inflammatory myopathies (IIM); polymyositis (PM) and dermatomyositis (DM) have serological profiles within the disease spectrum (7, 46, 112, 139). Myositis specific autoantibodies (MSAs) appear to be important prognostic markers and are associated with specific clinical features.

The aim of this study, described in this chapter, was to investigate the prevalence of recognised MSAs in the Royal National Hospital for Rheumatic Diseases (RNHRD) IIM cohort in comparison to previous published series (6, 46, 112). This study also highlights the clinical features associated with the different MSAs, in particular lung disease. The work described here has been presented at the *British Society of Rheumatology 2008* and the *American College of Rheumatology 2006*:

- Gunawardena H, Betteridge Z and McHugh NJ. The prevalence and clinical associations of myositis specific and associated autoantibodies in idiopathic inflammatory myopathy (abstract). *Rheumatology 2008*;47(S2):ii84.
- Gunawardena H, Robinson G, Betteridge Z, North J, Carmichael C, Foley NM, McHugh NJ. Interstitial pneumonia in polymyositis and dermatomyositis (abstract). *Arthritis Rheum. 2006*;54:S660.

##### 3.2 Patients and Methods

###### 3.2.1 Patients

Patients with a diagnosis of adult IIM (age of onset 18 years or over) were identified from the RNHRD CTD database and clinics. Cases of myositis / connective tissue disease (CTD) overlap were excluded. The clinical notes were reviewed to confirm the diagnosis of PM or DM based on the *Bohan and Peter* criteria (1, 2), and clinical data were retrospectively recorded using a standardised proforma (see Chapter 2, Appendix 1). Data recorded included details on initial presentation, creatinine kinase (CK) at disease onset and maximum CK during disease course. The degree of

cutaneous involvement and other features were noted including the presence of lung disease and cancer-associated myositis.

### **3.2.2 Assessment of interstitial pneumonia**

High resolution computerised tomography (HRCT) scans were re-reviewed under the guidance of Consultant Thoracic Radiologist (GR) and the subtype of interstitial pneumonia was categorised based on the American Thoracic Society / European Respiratory Society classification (32).

### **3.2.3 Serological Methods**

#### ***ANA-Immunofluorescence (IIF)***

IIF was previously performed by standard methods (PO and JD) using HEp-2 cells and fluorescein-labelled anti-human IgG immunoglobulin (Sigma, UK). The ANA IIF results were reviewed from the clinical notes and RNHRD / Bath Institute for Rheumatic Diseases serological database. The ANA titre and pattern was recorded.

#### ***Protein Radio-immunoprecipitation (IPP)***

Sera stored at -20°C were thawed at room temperature. IPP from K562 cell extracts was performed as previously described in Chapter 2, see Sections 2.3.2 and 2.3.3. Briefly, 10 µl sera was mixed with 2 mg protein-A-Sepharose beads (Sigma, UK) in IPP buffer (10 mM Tris-Cl pH 8.0, 500 mM NaCl, 0.1% v/v Igepal) at room temperature for 30 min. Beads were washed in IPP buffer prior to the addition of 120 µl [<sup>35</sup>S] methionine labelled K562 cell extract. Samples were mixed at 4°C for 2 hr. Beads were washed in IPP buffer followed by TBS buffer (10 mM Tris-Cl pH 7.4, 150 mM NaCl) before being resuspended in 50 µl SDS sample buffer (Sigma, UK). After heating, proteins were fractionated by 10% SDS-PAGE, enhanced, fixed and dried. Labelled proteins were analysed by autoradiography.

### **3.2.4 Ethical approval**

All patients gave fully informed written consent to participate and provide biological samples according to the Declaration of Helsinki under local ethical committee regulations.

## **3.3 Results**

Fifty patients with adult-onset IIM were identified from the RNHRD CTD database. The serological profiles and clinical associations are described below:

### 3.3.1 Serological profiles

Following IPP, autoantibody specificity was categorised into recognised MSAs, myositis-associated autoantibodies (MAAs), unidentified or negative. The frequency of recognised MSAs in this cohort of 50 patients was: anti-aminoacyl-tRNA synthetases (anti-ARS) 28%, anti-SRP 8% and anti-Mi-2 4%. Fourteen% of cases had MAAs (8% anti-PM-Scl, 4% anti-Ro, 2% anti-Ku). Twenty-two% had autoantibodies that appeared to recognise distinctive autoantigen bands (see Chapter 4). Twenty-four% of patients sera were either negative or precipitated non-specific bands by IPP. See Table 4 for summary and Figure 4.

### 3.3.2 Clinical associations

The main clinical features of patients with MSAs; anti-ARS, anti-SRP, anti-Mi-2 and MAAs (anti-PM-Scl, anti-Ku, anti-Ro) are described in Table 5. The clinical associations of the novel autoantibody specificities are described in Chapter 4. The main clinical features of anti-ARS positive patients (n=14) were arthritis 78.6%, mechanic's hands 57.1%, interstitial pneumonia 71.4% and Raynaud's phenomenon 71.4%, and 71.4% initially presented with arthritis or respiratory disease prior to myositis onset. Anti-SRP positive patients (n=4) were characterised by severe disease; including severe myopathy (median CK 11036 IU/L), with systemic features of weight loss (50%) or fever (50%), arthritis (75%) and Raynaud's phenomenon (75%) and cardiac involvement (50%). No anti-SRP patients had interstitial pneumonia. Muscle biopsies were available from two anti-SRP-positive patients demonstrating predominant muscle fibre necrosis, endomysial fibrosis and scanty inflammatory cell infiltrate. Both anti-Mi-2 positive cases had classic DM skin lesions and myositis but no other systemic organ involvement.

### 3.3.3 Subtypes of interstitial pneumonia

Ten (71.4%) anti-ARS-positive patients and two (29%) of MAA-positive patients had evidence of interstitial pneumonia on HRCT. No anti-Mi-2-positive, anti-SRP-positive or autoantibody-negative patients had evidence of interstitial pneumonia. As observed in previous studies (28-30, 47, 48), anti-ARS autoantibodies were a significant risk factor for interstitial pneumonia in comparison to other autoantibody subsets (anti-ARS-negative patients) ( $P_{uncorr}=0.02$ , odds ratio 2.4, 95% confidence interval 1.3-4.6). The interstitial pneumonia radiological subtypes are described in Table 6.

The main subtype of interstitial pneumonia observed in anti-ARS-positive cases was fibrotic non-specific interstitial pneumonia (NSIP), in 5 out of 10 patients. This HRCT appearance was defined by the degree of reticular changes, with predominately a subpleural and basal distribution. The fibrosing pattern was homogenous with absence of dense fibrosis i.e. no significant traction bronchiectasis or honeycombing. Three anti-ARS-positive patients (two anti-PL-7 and one with anti-PL-12 autoantibodies) had signs consistent with usual interstitial pneumonia (UIP) on HRCT, characterised by significant reticular changes including traction bronchiectasis and honeycombing. Four patients (two anti-Jo-1-positive and two anti-PM-Scl-positive cases) had HRCT features of limited NSIP i.e. predominately ground glass pattern, which is suggestive of fine fibrosis or alveolitis with a limited subpleural or basal distribution. One patient with limited NSIP had co-existent features of organising pneumonia (OP), characterised by patchy air-space consolidation, in the same distribution. See Figures 5-7 - HRCT images of corresponding interstitial pneumonia subtypes.

**Table 4: Serological profiles (RNHRD IIM study)**

Autoantibody	RNHRD cohort (n=50)	Frequency (%)
		Previous studies**
Anti-ARS		
Anti-Jo-1	16	20
Anti-PL-12	2	<5
Anti-PL-7	4	<10
Anti-OJ	2	<5
Anti-KS	0	<5
Anti-EJ	0	<10
Anti-Ha	0	<1
Anti-Zo*	4	0
Anti-SRP	8	5
Anti-Mi-2	4	<10
Novel autoantibodies*		
-p155/140 kDa target	14	21
-p140 kDa target	2	0
-p40/90 kDa target	6	0
Non-specific or negative	24	n/a
MAAs		
Anti-PM-Scl	8	<10
Anti-Ro	4	<10
Anti-Ku	2	<10

**\*Novel autoantibodies: anti-Zo, -p155/140, -p40/90 – see Chapter 4, p140 – see Chapter 7**

**\*\*References: (112, 143)**

**Table 5: Clinical characteristics of anti-ARS, anti-SRP, anti-Mi-2 and MAA-positive patients**

	Autoantibody				
	Anti-ARS (n=14)	Anti-SRP (n=4)	Anti-Mi-2 (n=2)	MAAs (n=7)	Negative* (n=12)
Female	79	50	100	71	100
Median age at onset (yrs)	51.5	48.5	62.5	36	62.5
Median CK at onset (IU/L)	1421	11036	4092	2347	655
Skin lesions					
Heliotrope	14	0	100	14	25
Gottron's	43	0	100	29	33
Periungal	36	25	0	29	25
V-sign	0	0	100	0	17
Shawl-sign	0	0	50	0	17
Mechanic's hands	57	0	0	43	8
Fever	57	50	0	29	8
Weight loss	36	100	0	57	8
Dysphagia	0	50	0	14	25
Raynaud's	71	75	0	71	25
Arthritis	79	75	0	57	25
Interstitial pneumonia	71	0	0	29	0
Cardiac	0	50	0	0	0
CAM	0	0	0	0	0

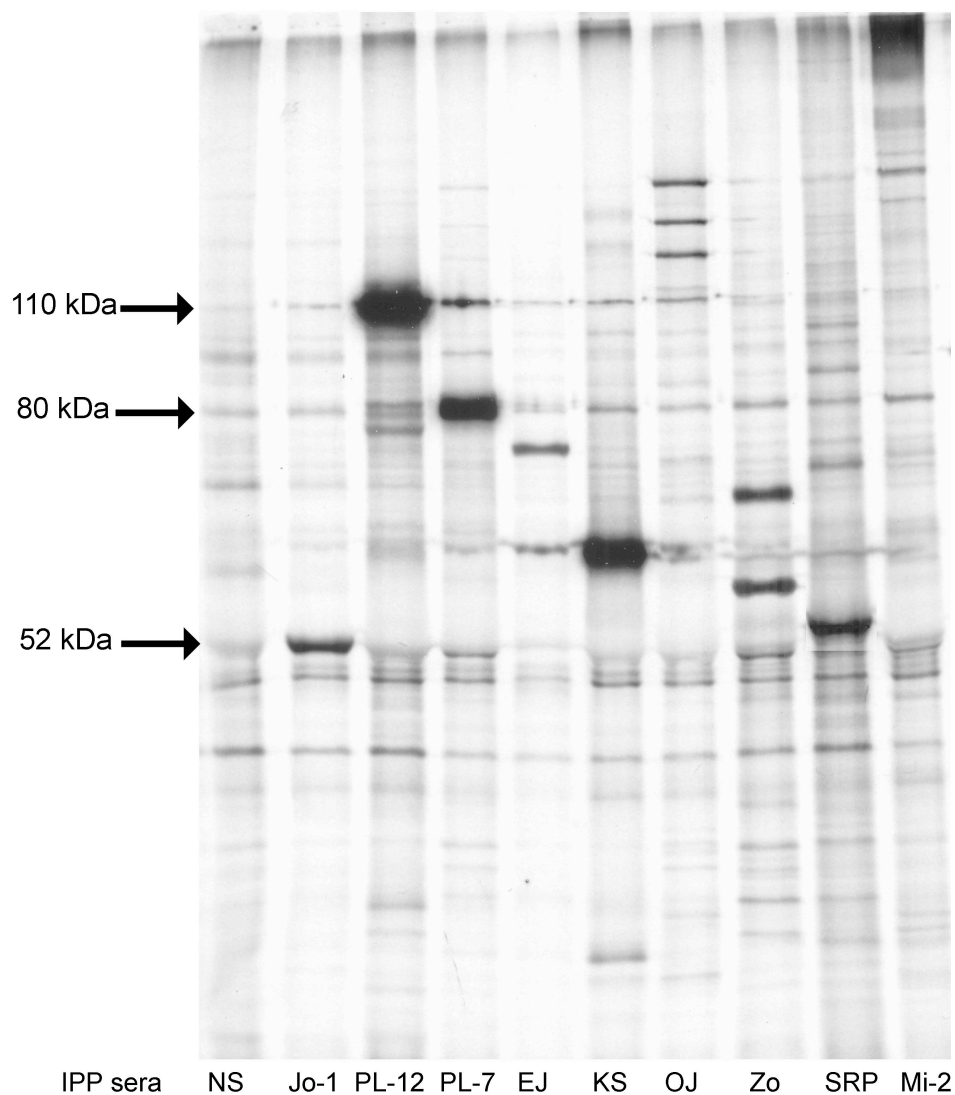
**\*Negative or non-specific unidentified bands on IPP**

**Percentages are shown unless otherwise stated**

**Figure 4: Immunoprecipitation of myositis-specific autoantigens**

Autoradiogram of 10% SDS PAGE of immunoprecipitates of [<sup>35</sup>S] labelled K562 cell extracts. The reference sera used for immunoprecipitation were; Lane 1: normal serum (QC4), Lane 2: anti-Jo-1 (QC78), Lane 3: anti-PL-12 (R12501), Lane 4: anti-PL-7 (R18134), Lane 5: anti-EJ (AOMIC MM182), Lane 6: anti-KS (AOMIC MM020), Lane 7: anti-OJ (AOMIC MM63), Lane 8: anti-Zo (R21845), Lane 9: anti-SRP (R7417), Lane 10: anti-Mi-2 (R18411).

Arrows indicate molecular weights (kDa) –see Chapter 2, Appendix, Table 4 for running molecular weights of 10% SDS=PAGE.



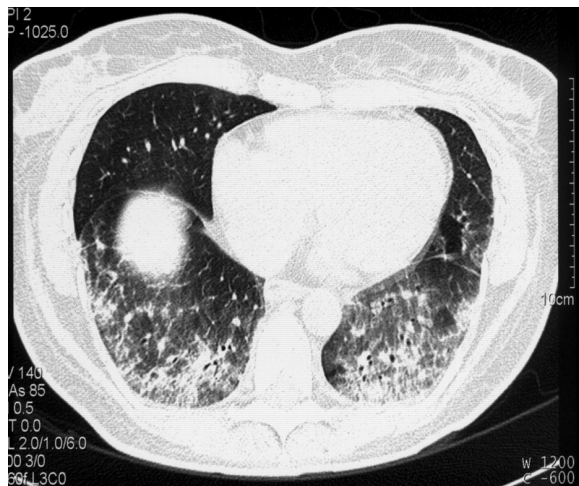
**Table 6: Subtypes of interstitial pneumonia in anti-ARS-positive and MAA-positive patients (RNHRD IIM study)**

<b>Patient</b>	<b>Autoantibody</b>	<b>HRCT IP subtype</b>
R16716	Anti-Jo-1	Fibrotic NSIP
R9435	Anti-Jo-1	Fibrotic NSIP
R13077	Anti-Jo-1	Limited NSIP
R5477	Anti-Jo-1	Fibrotic NSIP
R12707	Anti-Jo-1	Limited NSIP and OP
R18134	Anti-PL-7	UIP
R19896	Anti-PL-7	UIP
R12501	Anti-PL-12	UIP
R15222	Anti-Zo*	Fibrotic NSIP
R21845	Anti-Zo*	Fibrotic NSIP
R8125	Anti-PM-Scl	Limited NSIP
R10686	Anti-PM-Scl	Limited NSIP

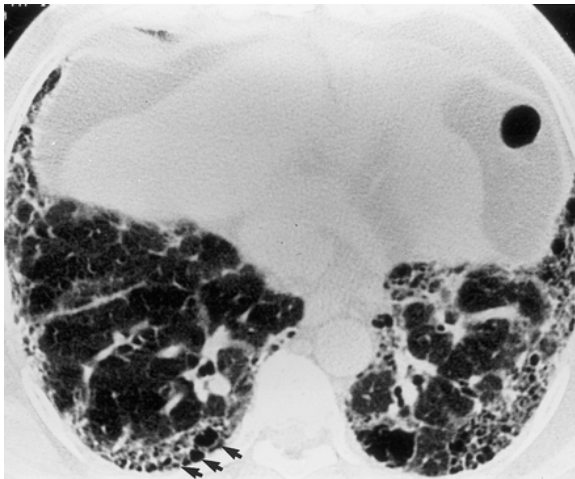
**\*See Chapter 4**



**Figure 5: Non-specific interstitial pneumonia**



**Figure 6: Usual interstitial pneumonia**



**Figure 7: Organising pneumonia**



### 3.4 Discussion

Historically, PM and DM has been defined based on broad pathological and clinical criteria (1, 2). There is now increasing evidence that these disorders characterised by a wide clinical spectrum can now be classified into more homogeneous clinico-serological syndromes (6, 46, 112, 139). Autoantibodies found exclusively in IIM, termed MSAs are directed against cytoplasmic or nuclear components involved in key regulatory processes such as protein synthesis, translocation and nuclear transcription, and have been the subject of major interest over the last few years. Unlike myositis-associated autoantibodies (MAAs) found in principally myositis-scleroderma overlap syndromes; MSAs are highly selective, mutually exclusive and are associated with characteristic clinical profiles. As described in Chapter 1; several myositis specific autoantigen systems have been described including the aminoacyl-tRNA synthetases (ARS) (111, 114-119), signal recognition particle (SRP) (121, 122) and Mi-2 (126).

Consistent with previous observations (112), in this study the most frequently detected MSA group were anti-ARS, with a lower frequency of anti-SRP and anti-Mi-2 autoantibodies. Using IPP, a number of novel autoantibody specificities were identified, which are described fully in Chapter 4.

The clinical manifestations of anti-ARS-positive patients were consistent with the anti-synthetase syndrome, with a high frequency of mechanic's hands, fever, arthritis, Raynaud's phenomenon and interstitial pneumonia. As previously described in anti-synthetase syndrome (42), the majority of patients presented with extra-muscular manifestations, in particular interstitial pneumonia, prior to their myositis onset. Clinically-amyopathic disease or milder muscle involvement is described especially in non-Jo-1 anti-ARS-positive cases (42, 144-148). In comparison to anti-ARS-negative cases, anti-ARS-positivity was a significant risk factor for the presence of interstitial pneumonia, as observed previously (6, 31, 42, 48, 144-147, 149-152). Consistent with previous findings (30, 38), NSIP was the frequent interstitial pneumonia subtype observed in anti-ARS-positive patients (including all anti-Jo-1-positives). Interestingly, anti-PL-7 or -12 positivity was associated with UIP, the radiological and histological pattern synonymous with idiopathic pulmonary fibrosis. Indeed, the predominant clinical manifestation in two of these patients was interstitial pneumonia, with one patient (anti-PL-7-positive) developing myositis approximately 10 years after the onset of lung disease and one

patient (anti-PL-12-positive) has never developed clinical myositis. In some cases, IPF may actually represent a *formes frustes* of autoimmune connective tissue disease, especially in patients with non-Jo-1 anti-ARS (150), which may not be tested for in routine clinical practice. In a study by *Matsushita et al* 25% of patients with idiopathic interstitial pneumonia were anti-ARS positive (152). At present, it remains unclear why Jo-1 is the most common autoantigenic target in ASS or why patients with anti-ARS autoantibodies have differences in their clinical presentation. In these patients, a high index of clinical suspicion including the recognition of other subtle clinical features, for example a history of arthralgia or fever, is required.

Similar to previous studies (122-124), all patients with anti-SRP autoantibodies had evidence of severe myopathy with markedly elevated CK at disease onset. In addition, patients had systemic features, cardiac involvement and arthritis but no significant skin disease, and none had evidence of IP. Therefore, identification of anti-SRP autoantibodies at disease onset has prognostic implications, identifying those patients at risk of severe muscle disease that may be refractory to standard treatments and thus require more intensive management. Anti-Mi-2 is a DM-specific autoantibody (6, 10, 125-128), and in this study, both patients with anti-Mi-2 autoantibodies had myositis associated with classic cutaneous DM lesions with no systemic features or internal organ involvement.

In summary, the MSAs: anti-ARS, -SRP, -Mi-2, identify IIM patients into three distinct clinical phenotypes. Rather than diagnose patients as purely DM or PM, if we adopt a serological classification to define patients into clinical syndromes this may help predict outcomes, in particular, identify those patients who are at risk of severe disease, which in turn may influence treatment strategies.

### 3.5 Appendix

#### I. SUBJECTS

Code	IIF pattern	IPP result	HRCT pattern
R12707	Coarse Cytoplasmic Speckle	Jo-1	Limited NSIP/OP
R16716	Cytoplasmic Speckle	Jo-1	Fibrotic NSIP
R13077	Discrete Cytoplasmic Speckle	Jo-1	Limited NSIP
R9435	FS	Jo-1	Fibrotic NSIP
R5477	Negative	Jo-1	Fibrotic NSIP
R4480	Discrete Speckles	Jo-1	n/a
R11739	Not available	Jo-1	n/a
R8452	FS and Discrete Cytoplasmic	Jo-1	n/a
R12501	Coarse Cytoplasmic Speckle	PL-12	UIP
R18134	Fine Cytoplasmic Speckle	PL-7	UIP
R19896	FSNS and Strong Fine Cytoplasmic Speckle	PL-7	UIP
R16682	Not available	OJ	n/a
R15222	Strong Cytoplasmic Speckle	Zo*	Fibrotic NSIP
R21845	Strong Cytoplasmic Speckle	Zo*	Fibrotic NSIP
R20227	FSNS and Strong Fine Cytoplasmic Speckle	SRP	n/a
R7417	Negative	SRP	n/a
R22034	Homogeneous with Fine Cytoplasmic Speckle	SRP	n/a
R12767	FSNS	SRP	n/a
R18411	Homogeneous	Mi-2	n/a
R16125	Homogeneous	Mi-2	n/a
R8125	Nucleolar	PM-Scl	Limited NSIP
R10686	Not available	PM-Scl	Limited NSIP
R17606	Homogeneous and nucleolar	PM-Scl	n/a
R14303	FS and nucleolar	PM-Scl	n/a
R9980	Negative	Ro	n/a
R15323	DCNS	Ro	n/a

Code	IIF pattern	IPP result	HRCT pattern
R21233	Homogeneous and Fine Cytoplasmic Speckle	Non-specific bands	n/a
R18506	FS and nucleolar	Non-specific bands	n/a
R13440	FSNS	Non-specific bands	n/a
R19013	FSNS	Ku	n/a
R13902	Homogeneous and Nucleolar	Non-specific bands	n/a
R13408	Negative	Negative	n/a
R7544	FS	Negative	n/a
R21308	FSNS	Negative	n/a
R10290	Negative	Negative	n/a
R14789	Homogeneous and Coarse Cytoplasmic Speckle	Negative	n/a
R9674	Homogeneous	Negative	n/a
R21140	DCNS	Negative	n/a
R19484	FSNS and Cytoplasmic Speckle	Negative	n/a
R20125	FSNS	155 and 140 kDa bands*	n/a
R9316	FSNS	155 and 140 kDa bands*	n/a
R7505	DCNS	155 and 140 kDa bands*	n/a
R16815	FSNS	155 and 140 kDa bands*	n/a
R8315	DCNS	155 and 140 kDa bands*	n/a
R13064	Homogeneous	155 and 140 kDa bands*	n/a
R11248	Homogeneous	155 and 140 kDa bands*	n/a
R18883	FS and Fine Cytoplasmic Speckle	140 kDa*	n/a
R16316	CS	40 and 90 kDa bands*	n/a
R11040	DCNS	40 and 90 kDa bands*	n/a
R20894	FSNS	40 and 90 kDa bands*	n/a

**\*See Chapter 4**

**N.A. – not applicable**

**ANA IIF patterns - abbreviations:**

C: coarse

D: diffuse

F: fine

S: speckle

NS: nucleolar sparing

## **II. Acknowledgements**

Mrs J Dunphy (JO) and Mrs P Owen (PO) previously performed ANA IIF on sera from IIM cases as part of the RNHRD (Bath Institute for Rheumatic Diseases) Diagnostic Laboratory Service. I would like to thank them for giving me permission to include this data in this study.

I would also like to acknowledge and thank Dr G Robinson (GR), Consultant Radiologist, Royal United Hospital, Bath for his supervision and teaching in the interpretation of HRCT scans.

I would like to thank all the patients who have attended the RNRHD CTD clinic and given their consent to participate in these studies.

## CHAPTER FOUR

### RESULTS

#### Identification and characterisation of novel autoantigen systems in adult-onset inflammatory myositis

##### 4.1 Introduction

To date, several myositis specific autoantibodies (MSAs) have been described including the anti-aminoacyl-tRNA synthetases (ARS) (6, 114-119), -signal recognition particle (SRP) (6, 123, 124) and -Mi-2 (6, 125-127) (as described in Chapter 1, see Table 2, for summary). Over the last few years, new MSAs have been described in adult IIM.

Sato *et al* have described an autoantibody termed anti-CADM-140 in a Japanese cohort of adult patients with clinically amyopathic dermatomyositis (CADM) (130). As defined by Gerami *et al*, patients with CADM have hallmark cutaneous features of DM but no clinical evidence of myositis (40). Moreover, CADM patients especially those of Asian origin, appear to have an increased risk of more severe interstitial pneumonia (31, 41, 153). In this study by Sato *et al*, anti-CADM-140 autoantibodies were detected in 8 out of 42 DM patients (19%) but not in patients with PM, other CTDs or control groups. All 8 patients with anti-CADM-140 autoantibodies were clinically amyopathic. Anti-CADM-140-positive sera demonstrated a granular or reticular cytoplasmic staining pattern on immunofluorescence (IIF) using HEp-2 cells. Following radio-immunoprecipitation (IPP) using <sup>35</sup>S-methionine-labelled K562 cell extract, anti-CADM serum recognised an ~140 kDa polypeptide, which was distinguishable from other known MSAs, and immunodepletion studies confirmed that this autoantibody recognises the same target. Of interest, the frequency of severe interstitial pneumonia was significantly increased in the anti-CADM-140-positive group compared to anti-CADM-140 negatives (50% versus 6%,  $p=0.008$ ) (130). Using a series of molecular techniques including a HeLa cell-derived complementary DNA library, the identity of the ~140 kDa protein has recently been identified as a RNA helicase encoded by melanoma differentiation-associated gene 5 (MDA5) (131). MDA5 is involved in innate immune responses against viruses (132, 154, 155). This finding suggests that a specific viral infection in genetically susceptible individuals may play a direct role in the pathogenesis of CADM and severe interstitial pneumonia, particularly as this MSA has been detected in Japanese patients, and confirmed in a further cohort recently (156).

A further novel MSA, termed anti-p155 or anti-p155/140 has been reported in adult DM. In the study by Targoff *et al*, following IPP using HeLa cell extract, anti-p155-positive sera recognised a distinct 155 kDa protein (along with a weaker 140 kDa protein in most cases) in 8 out of 39 DM patients (21%). Anti-p155 sera revealed a nuclear speckled pattern on IIF with HEp-2 cells (143). Kaji *et al* have demonstrated what is likely to be the same autoantibody, termed anti-p155/140, in 7 out of 52 DM patients (13%) using IIP with K562 cells (157). Both studies showed that anti-p155/140 autoantibodies also appear to identify a distinct clinical phenotype with more severe skin involvement. Moreover, the most striking feature is the clear association with malignancy and Chinoy *et al* have confirmed this. The authors showed that anti-p155/140 positivity confers a significant risk for cancer-associated-myositis with a high specificity, moderate sensitivity and high negative predictive value (158). In preliminary work, the identity of the p155 protein target is consistent with transcriptional intermediary factor 1-gamma (TIF-1- $\gamma$ ), a nuclear transcription protein involved in cellular differentiation (159).

This chapter describes data from the RNHRD IIM cohort on anti-p155/140 autoantibodies, and also work describing two novel MSAs (termed anti-Zo and anti-SAE). The work in this chapter has been published, as listed below:

- Betteridge Z, Gunawardena H, North J, Slinn J, McHugh N. Anti-synthetase syndrome: a new autoantibody to phenylalanyl transfer RNA synthetase (anti-Zo) associated with polymyositis and interstitial pneumonia. *Rheumatology*. 2007;46:1005-1008. (120)
- Betteridge Z, Gunawardena H, North J, Slinn J, McHugh N. Identification of a novel autoantibody in dermatomyositis directed against small ubiquitin-like modifier activating enzyme. *Arthritis Rheum*. 2007;56:3132-3137. (160)
- Gunawardena H, North J, Wedderburn LR, Davidson JE, Betteridge ZE, Dunphy J, Chinoy H, Cooper RG, McHugh NJ. Clinical associations of anti-p155/140 autoantibodies in adult and juvenile dermatomyositis. *Ann Rheum Dis*. 2007;66:S68. (161)
- Betteridge ZE\*, Gunawardena H\*, Chinoy H, North J, Ollier WER, Cooper RG, McHugh NJ. Clinical and HLA-class II haplotype associations of autoantibodies to small ubiquitin-like modifier enzyme, a dermatomyositis-specific autoantigen target, in UK adult-onset Caucasian myositis. *Ann Rheum Dis*. 2009;68:1621-5. (\*co-first author) (162)



## **4.2 Patients and Methods**

### **4.2.1 Patients**

Patients with a diagnosis of adult IIM (age of onset 18 years or over) were identified from the RNHRD CTD database and clinics. Cases of myositis / CTD overlap were excluded. The clinical notes were reviewed to confirm the diagnosis of PM or DM based on the *Bohan and Peter* criteria (1, 2), and clinical data were recorded using a standardised proforma (see Chapter 2, Appendix 1). Data recorded included details on initial presentation, creatinine kinase (CK) at disease onset and maximum CK during disease course. The degree of cutaneous involvement and other features were noted including the presence of lung disease and cancer-associated myositis. For the studies on anti-SAE autoantibodies (see Section 4.3.2) clinical data and serum samples were also available from the UK Adult Onset Myositis Immunogenetic Collaboration (AOMIC) (collaborators Dr Hector Chinoy and Dr Robert Cooper).

Sera from disease controls (150 with SSc / 40 with SLE) and 40 normal controls were also analysed (available from the RNHRD Bath Institute for Rheumatic Diseases Repository). All patients with SSc and SLE fulfilled recognised published criteria (11, 12).

### **4.2.2 Serological Methods**

#### ***ANA Immunofluorescence (IIF)***

IIF was previously performed by standard methods (PO and JD) using HEp-2 cells and fluorescein-labelled anti-human IgG immunoglobulin (Sigma, UK).

#### ***Protein Radio-immunoprecipitation (IPP)***

Sera stored at -20°C were thawed at room temperature. IPP from K562 cell extracts was performed as described in Chapter 2, see Sections 2.3.2 and 2.3.3. Briefly, 10 µl sera was mixed with 2 mg protein-A-Sepharose beads (Sigma, UK) in IPP buffer (10 mM Tris-Cl pH 8.0, 500 mM NaCl, 0.1% v/v Igepal) at room temperature for 30 min. Beads were washed in IPP buffer prior to the addition of 120 µl [<sup>35</sup>S] methionine labelled K562 cell extract. Samples were mixed at 4°C for 2 hr. Beads were washed in IPP buffer followed by TBS buffer (10 mM Tris-Cl pH 7.4, 150 mM NaCl) before being resuspended in 50 µl SDS sample buffer (Sigma, UK). After heating, proteins were fractionated by 10% SDS-PAGE, enhanced, fixed and dried. Labelled proteins were analysed by autoradiography.

### ***Immunodepletion experiments***

As described in Chapter 2, section 2.3.4: duplicate samples each containing 10 mg protein A Sepharose beads in 1 ml IPP buffer and 50 µl serum were mixed with end-over-end rotation at room temperature for 30 min. Beads were washed four times in 1 ml IPP buffer and 1 tube (A) was placed on ice whilst 150 µl [<sup>35</sup>S] methionine-labelled K562 cell extract and 350 µl IPP buffer was added to the remaining tube (B). Tube B was mixed at 4°C for 2 hr after which the supernatant was transferred to tube A, this was mixed at 4°C for a further 2 hr. The supernatant from tube A was then transferred to a fresh tube (C) and stored at -80°C. IPP using reference sera and either 150 µl control [<sup>35</sup>S] methionine-labelled cell extract or the immunodepleted supernatants (C) were completed as described for IPP using [<sup>35</sup>S] methionine.

### ***Protein isolation and mass spectrometry***

Autoantigen isolation and identification experiments were carried out by ZB (see Appendix 4.5 for details). Therefore, the methodology has not been described in Chapter 2, but is outlined in brief in this section. Forty µl of reference sera was mixed with 2 mg protein-A-Sepharose beads in 500 µl IPP buffer at room temperature for 30 min with end-over-end rotation. Beads were washed two times in 1 ml 0.2M triethanolamine pH 8.2 (Sigma, UK) and bound antibodies were crosslinked to the beads using 5mM bis-(sulphosuccinimidyl)-suberate (Perbio, UK) in 1 ml triethanolamine, mixing at room temperature for 30 min. The reaction was stopped with 1 ml 50mM Tris-Cl pH 7.5, mixing at room temperature for 15 min. The antibody coated Sepharose beads were washed three times in phosphate-buffered-saline and twice in IPP buffer prior to the addition of 1 ml K562 cell extract, corresponding to approximately  $1 \times 10^6$  cells. Samples were mixed end-over-end rotation at 4°C for 1 hr. The supernatant was removed and the beads were resuspended in a further 1 ml K562 cell extract and were mixed for 1 hr at 4°C. Beads were washed four times in IPP buffer and once in TBS before being resuspended in 80 µl SDS sample buffer. After heating (95°C for 4 min), proteins were fractionated by 10% SDS-PAGE. Gels were washed 3 x 5 min in pure water, stained for 60 min using Imperial Protein Stain (Perbio, UK) and destained overnight in pure water. Unique bands were removed to a 96-well plate.

### ***Mass Spectrometry (MS)***

Samples were prepared for MS at the University of the West of England using an Ettan robotic digester (GE Biosciences). Gel pieces were destained in 50%

Methanol / 50mM Ammonium Bicarbonate, dehydrated in 70% acetonitrile, air dried and digested overnight at room temperature with 20 ng/μl modified porcine trypsin (Promega) in 20mM ammonium bicarbonate. Peptides were extracted from gel pieces to a clean plate using 50% Acetonitrile / 0.1% TFA (trifluoroacetic acid) that was then dried down. The peptides were redissolved in 50% Acetonitrile / 0.1% TFA and mixed with an equal amount of 10 mg/ml alpha-cyano-4-hydroxycinnamic acid before being spotted on the MALDI target plate using an Ettan Robotic Spotter (GE Biosciences, UK). Peptide mass fingerprints were acquired using Waters Micromass Matrix-assisted laser desorption/ionisation-time of flight mass spectrometry (MALDI-TOF) with data acquisition and processing carried out using the MassLynx software. The database searching was performed by ProteinLynx software.

#### **4.2.3 Ethical approval**

All patients gave fully informed written consent to participate and provide biological samples according to the Declaration of Helsinki under local ethical committee regulations.

### **4.3 Results**

#### **4.3.1 Anti-Zo, a novel autoantibody in the anti-synthetase syndrome**

##### ***Identification of anti-Zo autoantibodies to phenylalanyl tRNA synthetase***

Prior to this study, seven anti-aminoacyl-tRNA (ARS) synthetase autoantibodies had been described (see Chapter 1, Section 1.6.4.1 and Table 1). As part of the work related to the study described in Chapter 3, it was noted that two IIM sera (R15222 and R21845) from the RNHRD cohort study had a strong discrete cytoplasmic speckle on IIF using HEp-2 cells, the pattern of which was suggestive of anti-ARS autoantibodies (see Figure 8). Protein IPP with the two prototype sera produced a novel pattern with two bands at approximately 60 and 70 kDa (see Figure 9A). The molecular weights of the bands immunoprecipitated did not match any of the known tRNA synthetases associated with myositis, including Jo-1, PL-7 and PL-12 (see Figure 9B) and as described for KS, EJ and OJ (see Chapter 3, Figure 4), implying that both index cases serum (R15222 and R21845) IPP pattern was due to a novel anti-ARS autoantibody. This pattern was not observed in any other disease or control groups.

In order to further characterise the anti-ARS, the corresponding autoantigen was purified and identified using SDS-PAGE and MALDI-TOF MS. Sera from either the index cases or patients with known autoantibodies to Jo-1, PL-7 or PL-12 were used

to immunoprecipitate autoantigens from a K562 cell extract. A Coomassie stained SDS-PAGE of the immunoprecipitates showed bands of expected molecular weight for the Jo-1, PL-7 and PL-12 controls (see Figure 9B) as well as 60 and 70 kDa bands from the index case. The bands were digested by trypsin, analysed by MS and SwissProt database matched. Matches required peptide coverage of over 20% and scores of approximately 12. All matches were repeated on at least two separate occasions for the controls and three separate occasions for the index case immunoprecipitates. Table 7 demonstrates the correct identification of Jo-1, PL-7 and PL-12 autoantigens in the control sera and the database matching of the autoantigens precipitated by the index case. On each of the three separate occasions, the 60 kDa band was matched to phenylalanyl tRNA synthetase alpha chain (57 kDa protein) and the 70 kDa band was matched to phenylalanyl tRNA synthetase beta chain (66 kDa protein).

### ***Clinical features of anti-Zo autoantibodies***

#### ***Index case 1 (R15222)***

A 49 year old woman presented with shortness of breath; six months later she developed proximal muscle weakness, Raynaud's phenomenon, puffy fingers, mechanic's hands and arthralgia. Clinical findings demonstrated a proximal myopathy with a creatinine kinase (CK) of 9533 IU/L. Muscle biopsy confirmed a necrotising myopathy with inflammatory cells. Pulmonary function tests revealed a restrictive pattern and high-resolution computerized tomography (HRCT) showed signs of fibrotic non-specific interstitial pneumonia (NSIP).

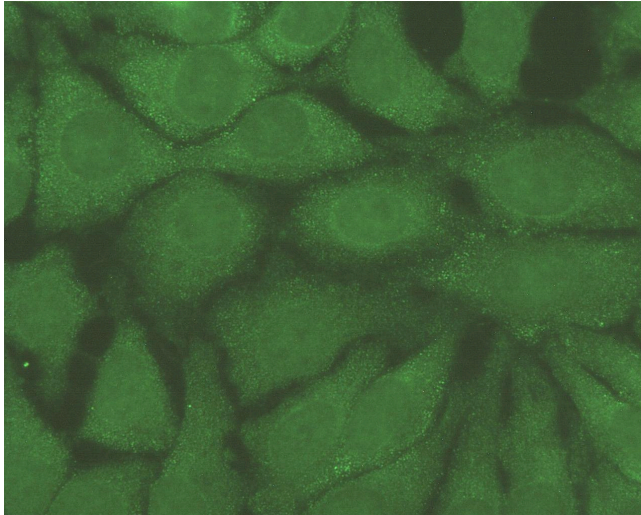
#### ***Index case 2 (R21845)***

A 34 year old man presented with shortness of breath on exertion. Over the next three months he developed Gottron's papules on the extensor aspects of his fingers, mechanic's hands, proximal myopathy with a CK of 4276 IU/L (muscle biopsy demonstrated a necrotising myopathy with inflammatory cell infiltrate), fever, Raynaud's and arthritis. HRCT thorax showed signs of fibrotic NSIP.

In summary, both patients with anti-Zo autoantibodies had typical clinical manifestations of the anti-synthetase syndrome.

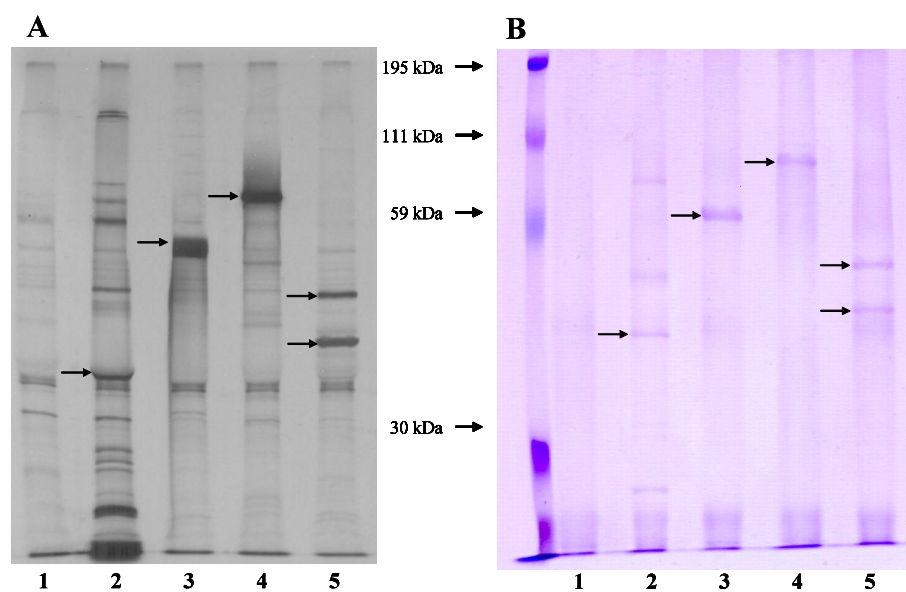
**Figure 8: ANA IIF – anti-Zo autoantibodies**

Indirect immunofluorescence staining of Hep-2 cells after incubation with index case 1 serum (R15222) and fluorescein-labelled anti-human immunoglobulin. The cytoplasmic speckle seen at 40x magnification is shown.



**Figure 9: IPP - anti-Zo autoantibodies**

**A** – Autoradiogram of 10% SDS-PAGE of immunoprecipitates using [<sup>35</sup>S] labelled K562 cell extracts. **B** – Coomassie stained 10% SDS-PAGE of immunoprecipitates using unlabelled K562 cell extract. Sera used for immunoprecipitation include; Lane 1; control serum; Lanes 2-5, anti-synthetase sera with autoantibodies to; Lane 2 – Jo-1; Lane 3 – PL-7; Lane 4 – PL-12 and Lane 5 – index case serum, R15222 (containing autoantibodies to phenylalanyl tRNA synthetase). Molecular weight markers are indicated on the left of panel B and bands corresponding to the tRNA synthetase targets are marked.



**Table 7** - Results of Matrix-assisted laser desorption/ionisation-time of flight mass spectrometry (MALDI-TOF MS) and SwissProt database matching. Comparison of peptide fragments of antigens precipitated by serum containing anti Jo-1, PL-7 and PL-12 along with peptides from the 60 kDa and 70 kDa bands precipitated by two index cases sera. Mass peptide fingerprints matched to histidyl-, threonyl- and alanyl-, phenylalanyl (alpha chain)- and phenylalanyl (beta chain)- tRNA synthetase respectively. Matches were deemed positive if the peptide coverage was over 20%, the same major theoretical and experimental peaks were present and the maximal MALDI-TOF score (using the software and database combination) was approximately 12.

Accession Number	Name	Description	<sup>1</sup> Mwt (Da)	<sup>2</sup> PI	Coverage	Score
P12081	SYH_Human	Histidyl tRNA synthetase (Jo-1)	57348	5.827	21.6%	11.7
P26639	SYTC_Human	Threonyl tRNA synthetase cytoplasmic (PL-7)	83381	6.592	32.0%	12.3
P49588	SYA_Human	Alanyl tRNA synthetase (PL-12)	106734	5.387	27.0%	11.5
Q9Y285	SYFB_Human	Phenylalanyl tRNA synthetase alpha chain	57396	7.881	29.2%	12.2
Q9NSD9	SYFB_Human	Phenylalanyl tRNA synthetase beta chain	66087	6.777	33.4%	12.3

<sup>1</sup>Mwt –theoretical molecular weight, <sup>2</sup>PI – theoretical isoelectric point.

### **4.3.2 Anti-SAE: autoantibodies targeting small ubiquitin-like modifier activating enzyme in adult dermatomyositis**

#### **4.3.2.1 Anti-SAE: autoantibodies targeting small ubiquitin-like modifier activating enzyme in adult dermatomyositis (RNHRD study)**

##### ***Identification of anti-SAE autoantibodies in index cases***

The RNHRD IIM study described in Chapter 3 identified two DM patients with the same previously unidentified autoantibody pattern (R16316 and R11040). Sera from these patients gave a speckled nucleolar sparing staining pattern on IIF (R16316 serum also showed a fine cytoplasmic speckle) (data not shown) and were found to immunoprecipitate proteins of approximately 40 kDa and 90 kDa (see Figure 10) that did not correspond to any known autoantigen profile. This pattern was not observed in any other disease or control groups.

##### ***Autoantigen identification***

Immunodepletion studies were completed to demonstrate that the IPP pattern seen with R16316 and R11040 were due to the precipitation of the same autoantigens. Cell extracts were depleted of antigen targets using normal, R16316 and R11040 sera before being used in further IPP using R16316 and R11040 sera. When the cell extracts were pre-depleted with normal serum, the 40 and 90 kDa antigens were still visible after IPP with each of the patient's sera (see Figure 10, Lanes 2 and 6). However, after pre-depletion with either patient's sera and removal of the corresponding targets, the 40 and 90 kDa bands were absent after subsequent IPP by either patient's sera (see Figure 10, Lanes 3-4 and 7-8). These results therefore provided good evidence that the serum from R16316 and R11040 both contained the same autoantibody specificity.

##### ***Identification of the SAE antigens***

Non-radiolabelled IPP and Coomassie stained SDS-PAGE demonstrated the presence of 40 kDa and 90 kDa bands using both R16316 and R11040 serum (see Figure 11). MALDI-TOF MS and Swiss Prot analysis of the peptide fingerprints from these bands corresponded to SUMO 1 Activating Enzyme A subunit (SAE1) (38 kDa protein) for the 40 kDa band and SUMO 1 activating enzyme B subunit (SAE2) (71 kDa protein) for the 90 kDa band (see Table 8).



### ***Clinical features of anti-SAE autoantibodies in index cases***

#### ***Index Case 1 (R16316)***

A 52 year old lady presented with a three month history of DM rash, including periungal erythema, V-sign rash and heliotrope rash. She had no muscle symptoms with a normal CK and electromyography (EMG). Six months later she developed proximal myopathy (CK 797 IU/L), dysphagia, and worsening skin disease with the Shawl-sign rash. Muscle biopsy confirmed necrotising myopathy with persistent inflammatory change. Screening investigations showed no sign of an underlying malignancy. HRCT showed limited peripheral lung involvement consistent with limited mild NSIP.

#### ***Index Case 2 (R11040)***

A 62 year old man presented with clinical features of cutaneous DM, including a heliotrope rash and Gottron's papules. Six months later he developed proximal myositis with pharyngeal involvement. CK was 1086 IU/L and the diagnosis was confirmed following EMG study. Further investigations excluded an underlying malignancy but did show changes consistent with peripheral limited mild NSIP.

In summary, the results from this study suggest that anti-SAE autoantibodies may form a further serological subset in adult IIM with similar clinical manifestations. A further patient (R20894) was subsequently identified as anti-SAE-positive, with this finding included in the anti-SAE UK AOMIC study described below.

**Table 8** – Results of MALDI-TOF MS and SwissProt database matching. Comparison of mass peptide fragments of antigens precipitated by patient 1 and 2 sera to small ubiquitin like 1 activating enzyme (SAE1) and (SAE2). Results from a total of three separate occasions (with two matches for R16316 and one match for R11040) are shown. Matches were deemed positive if the peptide coverage was over 20 percent and the same major theoretical and experimental peaks were present.

Accession Number	Name	Description	<sup>1</sup> Mwt (Da)	<sup>2</sup> PI	Coverage
Q9UBE0	ULE1A_ Human	Ubiquitin like 1 activating enzyme	38426	5.23	47.1%
		E1A SUMO1 (activating enzyme			46.2%
		subunit 1)			41.0%
Q9UBT2	ULE1B_ Human	Ubiquitin like 1 activating enzyme	71179	5.19	25.8%
		E1B SUMO1 (activating enzyme			37.8%
		subunit 2)			29.4%

<sup>1</sup>Mwt –theoretical molecular weight, <sup>2</sup>PI – therotical isoelectric point.

#### **4.3.2.2 Anti-SAE autoantibodies in UK AOMIC cohort**

The AOMIC registry has recruited UK Caucasian patients with adult-onset myositis, 18 years or over at disease onset, from hospitals around the UK. All patients had probable or definite myositis according to the *Bohan and Peter* diagnostic criteria (1, 2). A standardised proforma, including demographic and clinical data, was used throughout (see Chapter 2, Appendix). Cancer associated myositis defined as cancer occurring within 3 years of diagnosing myositis (as per the modified *Bohan and Peter* classification used in a previous study (9)) was confirmed by the local clinician caring for individual patients. Patients were investigated for interstitial pneumonia based on clinical symptoms and pulmonary function testing and where indicated HRCT thorax. Clinical data and serum samples from 266 myositis patients were available for analysis for this study. The median age of disease onset was 50 years (IQR 37-61) and 195 (73%) cases were female. One hundred and thirty one cases had DM, 124 PM and 11 myositis / CTD overlap. Sera and DNA samples were taken at the time of diagnosis and stored at -20°C until required.

#### ***Identification of anti-SAE autoantibodies in UK AOMIC cohort***

Sera from 266 myositis patients were screened for anti-SAE autoantibodies. Excluding the reference cases (R16316 and R11040) (as described in Section 4.3.2.1), nine further patients were identified as having 40 kDa and 90 kDa bands on SDS-PAGE, previously described as corresponding to SAE1 and SAE2 respectively (see Section 4.3.2.1 and Reference (160)). The presence of anti-SAE autoantibodies in these sera was confirmed by immunodepletion experiments. Immunoprecipitations on the 40kDa and 90 kDa positive sera were completed using cell extracts depleted with either the index case serum (anti-SAE), or with healthy control serum. All sera were positive for the 40 kDa and 90 kDa bands using healthy control depleted cell extract but were negative for the 40 kDa and 90 kDa bands with the index case (anti-SAE) depleted extract (see Figure 12). These results provide confirmation that the serum from the index cases and the nine additional patients contain the same autoantibody specificity, anti-SAE. Anti-SAE autoantibodies were not detected in any of the disease or normal control groups.

None of the 11 anti-SAE positive sera were found to co-immunoprecipitate any other recognised MSAs, although 5 patients did co-immunoprecipitate additional unidentified non-specific bands on IPP. IIF on anti-SAE positive sera demonstrated a fine speckled nucleolar-sparing pattern (titres ranging from 1/160 to 1/2560) in ten

patients with two sera also showing a fine cytoplasmic speckle (1/40 titre) and a homogeneous pattern (1/640 titre) in one case.

#### ***Clinical features of anti-SAE autoantibodies in UK AOMIC cohort***

The overall frequency of anti-SAE autoantibody positive cases in the myositis cohort was 4%. Anti-SAE was found exclusively in DM patients (n=131), with a frequency of 8%. The detailed clinical features of the 11 anti-SAE positive patients are described in Table 9. Seven (64%) were female and the mean age was 62 years (IQR 54-68). Nine patients (82%) had a heliotrope rash, 9 (82%) patients had Gottron's lesions on the fingers and 7 (64%) had Gottron's over the large joints (elbow or knee). Eight out of 8 patients (100%) (data not available on 3 patients) had periungual changes. Seven out of 9 patients (78%) (data not available on 2 patients) had dysphagia, 3 of which had profound swallowing dysfunction, which required enteral feeding. Nine out of 11 patients (82%) had systemic features as defined by fever, weight loss and raised inflammatory markers. Two patients (18%) had cancer-associated myositis. Two patients (18%) had radiological evidence of ILD (limited non-specific interstitial pneumonia with less than 10% lung involvement on HRCT scored by a thoracic radiologist) but neither of these 2 patients had respiratory symptoms. Seven anti-SAE positive cases (78%) (data not available on 2 patients) presented with cutaneous DM only (no weakness and normal CK), but then went onto develop frank myositis. The median disease duration between the initial presentation of skin disease and myositis onset was 3 months (IQR 1, 6). After multiple corrections were applied, no significant clinical associations were observed when comparing anti-SAE positive patients with anti-SAE negative DM patients except for a higher frequency of periungual changes in the anti-SAE positive group ( $P_{corr}=0.03$ ). In addition, data on the type of presentation i.e. skin disease first or skin / muscle disease at onset was not available on the anti-SAE negative DM group to make this comparison. The data is summarised in Table 10. This study has confirmed that SAE is an important autoantigen target in DM with specific clinical associations.

**Table 9: Clinical features of patients with anti-SAE autoantibodies in UK AOMIC cohort**

	<b>Patient</b>										
	1	2	3	4	5	6	7	8	9	10	11
<b>Age at onset</b>	51	62	75	46	59	75	63	52	56	78	66
<b>Sex</b>	F	M	F	M	F	F	M	F	M	F	F
<b>Heliotrope</b>	Yes	Yes	Yes	Yes	Yes	Yes	Yes	No	Yes	Yes	No
<b>Gottron's</b>											
<b>Fingers*</b>	No	Yes	Yes	Yes	No	Yes	Yes	Yes	Yes	Yes	Yes
<b>Other**</b>	No	No	Yes	Yes	No	No	Yes	Yes	Yes	Yes	Yes
<b>Periungal</b>	Yes	Yes	Yes	Yes	Yes	Yes	Yes	Yes	~	~	~
<b>V-sign</b>	Yes	Yes	No	Yes	No	No	No	~	~	~	~
<b>Shawl-sign</b>	Yes	Yes	No	Yes	No	No	No	~	~	~	~
<b>MH</b>	No	No	No	No	No	No	No	~	~	~	~
<b>Dysphagia</b>	Yes	Yes	Yes	Yes	No	No	Yes	Yes	~	Yes	~
<b>Systemic</b>	Yes	Yes	Yes	Yes	No	No	Yes	Yes	Yes	Yes	Yes
<b>Weakness<sup>#</sup></b>	Yes	Yes	Yes	Yes	Yes	Yes	Yes	Yes	Yes	Yes	Yes
<b>Raised CK</b>	Yes	Yes	Yes	Yes	Yes	No	Yes	Yes	Yes	No	Yes
<b>ILD</b>	Yes	Yes	No	No	No	No	No	No	No	No	No
<b>CAM</b>	No	No	No	No	No	No	Yes	Yes	No	No	No
<b>Arthritis</b>	No	No	No	No	No	No	Yes	No	No	No	Yes
<b>Presentation (months)<sup>@</sup></b>	S (6)	S (6)	S (6)	S (3)	S/M	S (2)	S (3)	S (2)	~	S/M	~

**Key to Table 9:**

\*Gottron's lesions over the extensor aspects of fingers

\*\*Gottron's lesions over the extensor aspects of elbows or knees

~Clinical information not available

#Weakness – objective signs defined by manual muscle testing

MH (mechanic's hands)

ILD (interstitial lung disease) – patients asymptomatic / limited mild non-specific interstitial pneumonia on high resolution CT scan

CAM (cancer-associated myositis)

@S=presented skin disease first (number of months before myositis onset), S / M = presented with skin and muscle disease

Cases 1 and 2 (index cases R16316 and R11040 – RNHRD study)

**Table 10: Comparison of selected clinical features in anti-SAE positive vs. anti-SAE negative dermatomyositis patients\* (UK AOMIC cohort study)**

	<b>Anti-SAE positive (n=11)**</b>	<b>Anti-SAE negative (n=120)**</b>
<b>Females</b>	64	74
<b>Heliotrope</b>	82	71
<b>Gottron's papules (fingers)</b>	82	63
<b>Gottron's lesions (other)</b>	64	48
<b>Periungal erythema</b>	100 #	46
<b>V-sign rash</b>	43	35
<b>Shawl-sign rash</b>	43	37
<b>Dysphagia</b>	78	43
<b>Systemic</b>	82	54
<b>Weakness</b>	100	89
<b>Elevated CK</b>	82	85
<b>Interstitial pneumonia</b>	18	18
<b>CAM</b>	18	16
<b>Arthritis</b>	18	22

\*Values are the percentage of patients who had the clinical feature. \*\*Not all patients had data on each clinical feature. CAM: cancer-associated myositis.

# $P_{\text{corr}}=0.03$  compared to anti-SAE negative DM patients.

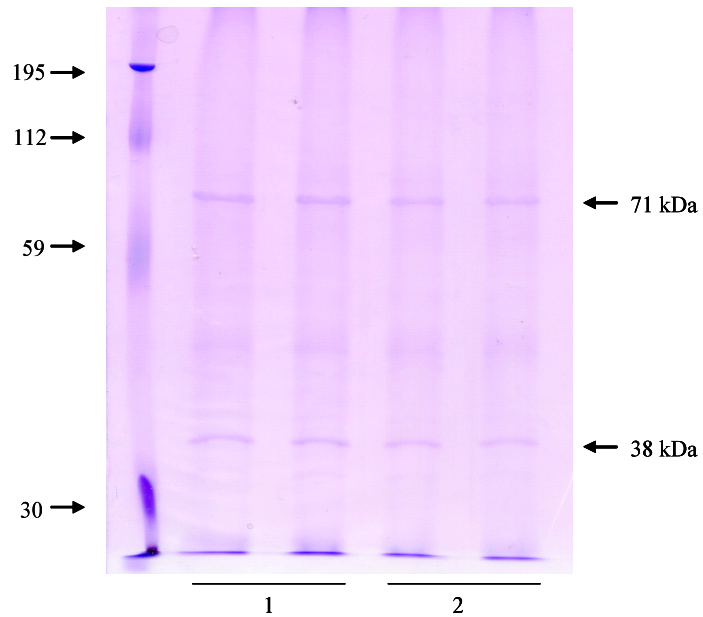
IPP of antigens recognised by patients R16316 and R11040. Autoradiogram of 10% SDS-PAGE of immunoprecipitates using either R16316 and R11040 serum with control [<sup>35</sup>S] methionine-labelled cell extract or [<sup>35</sup>S] methionine-labelled cell extract depleted with either normal sera (NS), patient R16316 sera (1) or patient R11040 sera (2). The bands corresponding to SAE1 (40 kDa band) and SAE2 (90 kDa band) are indicated.





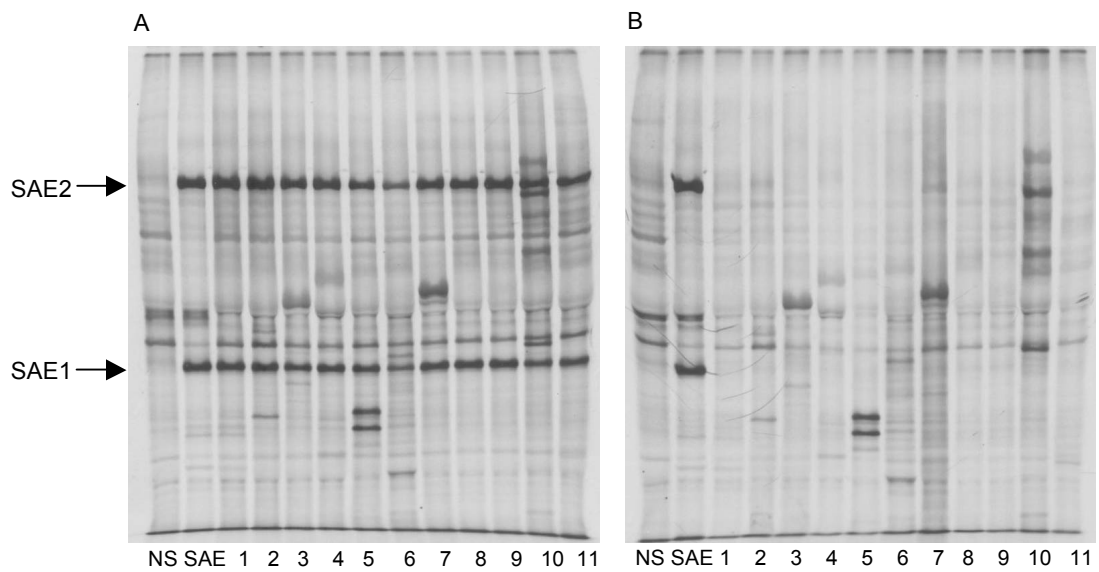
**Figure 11: Coomassie stained IPP of proteins using anti-SAE antibodies**

Coomassie stained 10% SDS-PAGE of immunoprecipitates using either patient 1 (R16316) or 2 (R11040) crosslinked serum precipitated with unlabelled K562 cell extract. The molecular weight markers are indicated in kDa on the left and the bands corresponding to SAE1 (40 kDa band) and SAE2 (90 kDa band) are indicated on the right.



### Figure 12: SAE Immunodepletion

Autoradiogram of 10% SDS-PAGE of immunoprecipitates using anti-SAE positive patient serum and either normal serum depleted [ $^{35}$ S] methionine – labelled cell extract (A) or SAE depleted [ $^{35}$ S] methionine – labelled cell extract (B) NS – IPP of normal serum and undepleted [ $^{35}$ S] methionine – labelled cell extract, SAE – IPP using anti-SAE positive serum and undepleted  $^{35}$ S methionine. Bands corresponding to SAE1 (40 kDa band) and SAE2 (90 kDa band) are indicated.



### **4.3.3 Anti-p155/140: autoantibodies targeting 155/140 kDa doublet protein in adult dermatomyositis**

#### ***Identification of anti-p155/140 autoantibodies***

The RNHRD IIM study described in Chapter 3 identified seven DM patients with the same previously unidentified autoantibody pattern and similar clinical features:

R7505	R13064
R8315	R16815
R9316	R 20125
R11248	

Following IPP testing, sera from the above cases of adult DM recognised two distinct proteins forming a doublet with molecular weights of 155 kDa and 140 kDa (see Figure 13). In each case a weak, non-specific nuclear pattern was detected on IIF (data not shown).

Immunodepletion studies were undertaken in order to ascertain whether the IPP pattern seen in the adult DM cases was due to precipitation of the same autoantigens. Immunoprecipitations on the different anti-p155/140 kDa positive sera were completed using cell extracts depleted with a reference index case anti-p155/140 serum (R7505), or with healthy control serum. All sera were positive for the p155/140 kDa bands using healthy control depleted cell extract but were negative for the p15/140 kDa bands with the index case (R7505) depleted extract (see Figure 14). These results provide confirmation that the serum from the index case and the six additional patients contain the same autoantibody specificity, anti-p155/140. None of the seven anti-p155/140 positive sera were found to co-immunoprecipitate any other recognised MSAs.

#### ***Frequency of anti-p155/140 autoantibodies***

A total of seven out of 50 (14%) adult IIM patients were positive for autoantibodies to the p155/140 kDa doublet. Anti-p155/140 was only detected in adult DM cases with a frequency of 26.9%. Anti-p155/140 autoantibodies were not found in any of the disease or normal control groups.

#### ***Clinical features of anti-p155/140-positive patients***

Findings are summarised in Table 11 and Table 12. Four adult DM patients with anti-p155/140 had a history of malignancy, which was diagnosed either at onset or

within three months of their DM. The remaining three anti-p155/140 patients had clinically-amyopathic DM (CADM) based on clinical findings and normal creatinine kinase levels since diagnosis, and no history of malignancy (median seven years from diagnosis to time of study).

In comparison to anti-p155/140 negative adult DM patients, anti-p155/140-positive patients had a higher frequency of cutaneous lesions; including V-sign rash, Shawl sign rash, heliotrope rash and Gottron's lesions. In addition, systemic manifestations including dysphagia were more frequent. No anti-p155/140-positive patients had interstitial pneumonia.

There was a significant association with anti-p155/140-positivity and malignancy ( $P_{corr}=0.03$ , odds ratio 7.3). The types of cancer seen in this group of patients were varied but all patients had evidence of solid organ malignancy.

**Table 11: Clinical features of anti-p155/140-positive adult DM vs. anti-p155/140-negative adult DM <sup>†</sup>**

	<b>Anti-p155/140 (adult DM)</b>	
	<b>Positive (n=7)</b>	<b>Negative (n=19)</b>
<b>Median age at diagnosis (yrs)</b>	54 (33, 71)	51 (49, 58)
<b>Females</b>	71.4	89.5
<b>Heliotrope rash</b>	85.7	63.2
<b>Gotttron's papules</b>	100	84.2
<b>Periungal erythema</b>	71.4	63.2
<b>V-sign rash</b>	71.4	26.3
<b>Shawl-sign rash</b>	85.7	26.3
<b>Dysphagia</b>	57.1	21.0
<b>Weight loss</b>	28.6	26.3
<b>Weakness</b>	57.1	73.7
<b>Elevated CK</b>	57.1	78.9
<b>Interstitial pneumonia</b>	0	31.6
<b>CAM</b>	57.1 #	0
<b>Arthritis</b>	28.6	36.8
<b>Calcinosis</b>	14.3	0
<b>Mechanics hands</b>	0	15.8

<sup>†</sup>Values are in percentages unless otherwise indicated

CK: creatinine kinase

CAM: cancer-associated myositis

# $P_{\text{corr}}=0.03$  compared to anti-p155/140 negative DM patients ( $P=0.002$ ), odds ratio 7.3, 95% CI 2.56-20.99)

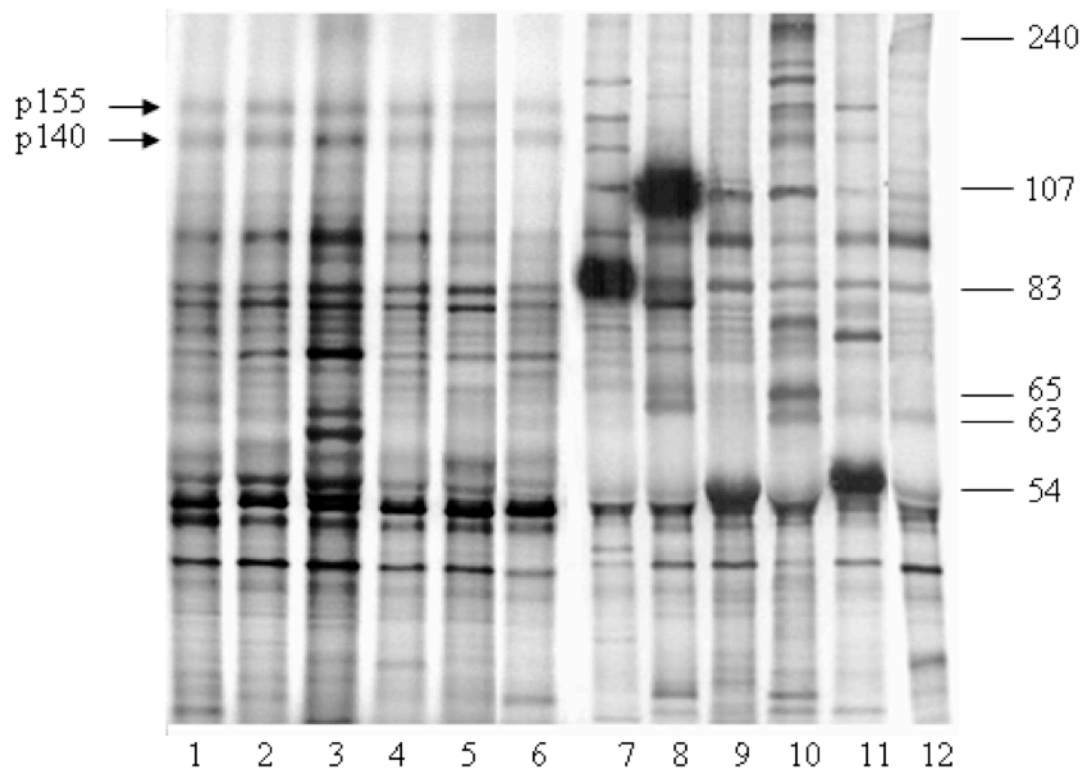
**Table 12: Anti-p155/140-positive patients (types of malignancy)**

<b>Patient</b>	<b>Diagnosis</b>	<b>Age at onset</b>	<b>Malignancy</b>
<b>1</b>	DM	82	Small cell carcinoma adrenal metastasis
<b>2</b>	DM	71	Bone and liver metastasis
<b>3</b>	DM	50	Breast carcinoma
<b>4</b>	DM	77	Gastric carcinoma
<b>5</b>	CADM	28	No
<b>6</b>	CADM	24	No
<b>7</b>	CADM	56	No

CADM: clinically-amyopathic dermatomyositis

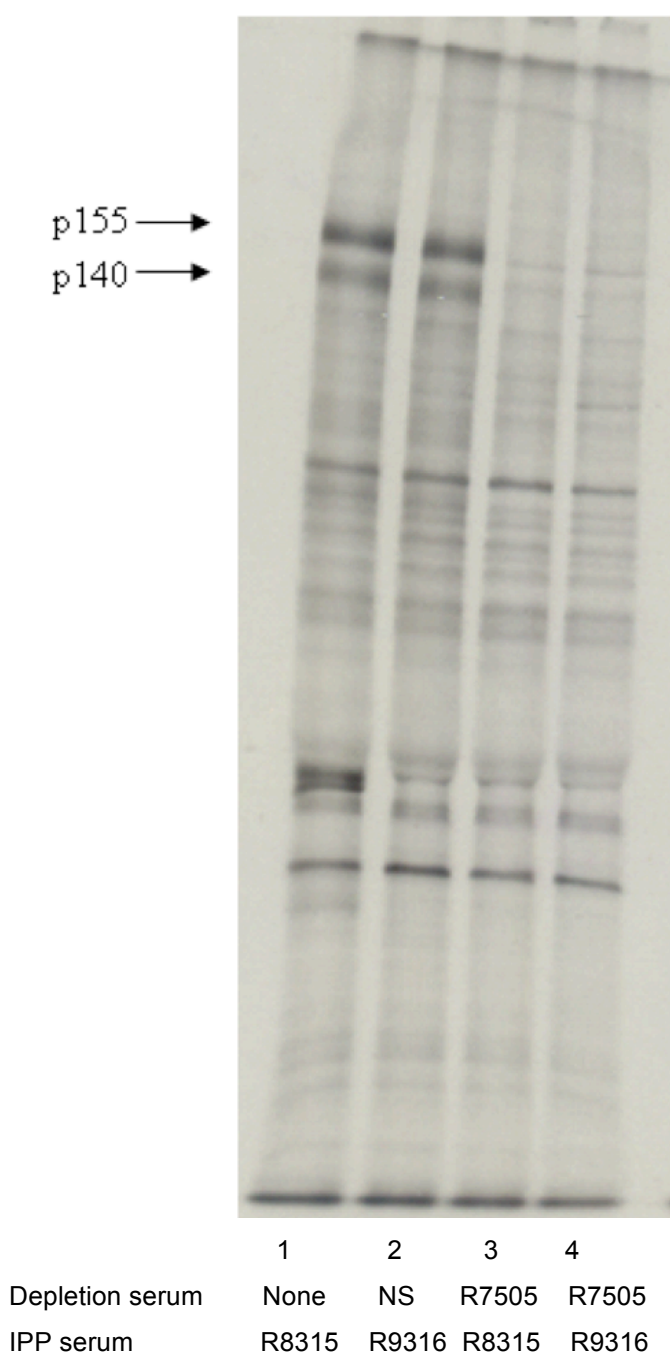
### Figure 13: IPP anti-p155/140 autoantibodies

Immunoprecipitation of p155/140 kDa autoantigens and other selected MSAs. 10% SDS PAGE of immunoprecipitates of [<sup>35</sup>S] labelled K562 cell extracts. Sera used for immunoprecipitation include Lanes 1-6; adult anti-p155/140 positive serum, Lane 7; threonyl tRNA synthetase, Lane 8; alanyl tRNA synthetase, Lane 9; histidyl tRNA synthetase, Lane 10; Mi-2, Lane 11; signal recognition particle and Lane 12; normal serum. Positions of the p155 and p140 antigens are indicated on the left. Positions of the Mi-2 bands at 240, 65 and 63 kDa are indicated. The 54, 83 and 108 kDa molecular weight markers correspond to signal recognition particle, threonyl tRNA synthetase and alanyl tRNA synthetase respectively.



**Figure 14: Immunodepletion experiments (anti-p155/140)**

Immunodepletion experiments - Autoradiogram of 10% SDS-PAGE of immunoprecipitates using adult positive anti-p155/140 serum: R7505, R8315 and R9316. Immunoprecipitation was performed with control [ $^{35}$ S] methionine-labelled cell extract or [ $^{35}$ S] methionine-labelled cell extract depleted with either normal serum (NS) or adult positive anti-p155/140 serum (R7505). The bands corresponding to the p155 and p140 autoantigens are indicated.





## **4.4 Discussion**

This chapter outlines work that characterises three new MSAs in adult IIM. The first is a new anti-ARS (-Zo) that targets phenylalanyl tRNA synthetase, which is the eighth autoantigenic ARS to be identified. The second is an autoantibody that targets SAE, an enzyme involved in post-translational modification, a completely novel system in adult DM. Finally, the clinical specificity of anti-p155/140 autoantibodies are described in patients recruited to the RNHRD IIM cohort study.

### **4.4.1 Anti-Zo; autoantibodies against phenylalanyl tRNA-synthetase**

The aminoacyl tRNA-synthetases are a distinct group of enzymes that catalyse the ATP-dependent binding of an amino acid to its cognate tRNA during protein synthesis. ARS are functionally related enzymes; each amino acid has its own corresponding tRNA-synthetase, except glutamic acid and proline, which have both been shown to be catalysed by a single polypeptide encoding the 'gluprolyl' tRNA synthetase (163). Previously, seven anti-ARS have been described (see Chapters 1 and 3) (111, 114-119). ARS belong to one of two classes depending on the amino acid they are responsible for. Both classes are multi-domain proteins with a catalytic domain and an anti-codon binding domain (ensures binding of the correct tRNA to the protein). Five ARS; histidyl (Jo-1), threonyl (PL-7), alanyl (PL-12), asparaginyl (KS) and glycyl (EJ) are class II synthetases found free in the cell cytoplasm. They have three highly conserved motifs, are generally dimeric or tetrameric, and attach their amino acid to the 3' hydroxy (OH) of their tRNA. The other two autoantigenic ARS: isoleucyl (OJ) and tyrosyl (Ha) are class I enzymes with two highly conserved motifs, are generally monomeric, and attach the carboxyl of their target amino acid to the 2' OH of adenosine 76 in the tRNA molecule. Anti-OJ autoantibodies are primarily directed against isoleucyl-tRNA synthetase but can react against multiple synthetases as part of a class I multi-enzyme complex including leucyl and glutaminyl-tRNA (117). Prior to this study; several ARS have not been shown to be autoantigenic (see Table 13). Either these synthetases are not targeted as part of an autoimmune response or autoantibodies directed against them are extremely rare. It is also feasible that it has not been possible to identify autoantibodies to the remaining synthetases using conventional diagnostic techniques.

Phenylalanyl tRNA synthetase is the eighth autoantigen associated with anti-synthetase syndrome to be identified. Similar to the majority of autoantigenic ARS, phenylalanyl is a class II synthetase but it aminoacylates at the 2' OH of the same adenosine, rather than at 3' OH.

Manifestations of anti-synthetase syndrome are described in Chapters 1 and 3. It is noteworthy that the two index cases identified as having anti-Zo autoantibodies had clinical features consistent with this syndrome. In particular, as well as myositis, both patients developed NSIP, the presence of which influenced treatment. As highlighted in Chapters 1 and 3, a major and sometimes predominant manifestation of anti-synthetase syndrome is interstitial pneumonia, which can be the major factor in terms of prognosis (42, 48). The different manifestations are not universal between patients and it is possible that each anti-ARS may define more specific clinical subsets within the syndrome itself. This is perhaps most notable in patients with non-Jo-1-anti-ARS (-KS, -PL-12, -PL-7) with lung disease in the absence of clinically apparent myositis (42, 48, 118, 144, 145). In a study by Yoshifuji H *et al*, the frequency of interstitial lung involvement in their patients with anti-synthetase syndrome was as high as 95%. Interestingly in terms of interstitial pneumonia being the first disease manifestation; anti-ARS autoantibodies were detected in 77% of this subgroup of patients with lung disease preceding myositis by a median of 11 months. However, in patients who developed myositis first, anti-ARS were only found in 20%. This study also looked at patients with interstitial pneumonia *sine* myositis; defined by anti-ARS, lung disease, no muscle disease either clinically or biochemically and no cutaneous IIM signs. All patients in this subgroup had non-Jo-1-anti-ARS (48). Similarly, the index anti-Zo-positive cases in this study initially presented with respiratory symptoms; the onset of myositis with other characteristic features occurring later. It is possible that non-Jo-1-anti-ARS may identify patients at one end of the spectrum. The question remains do anti-ARS have a predictive value in the context of idiopathic interstitial pneumonia i.e. do a proportion of these patients actually have a “*forme fruste*” of autoimmune CTD, as suggested in previous studies (48, 150). Detection of anti-ARS autoantibodies in these patients may influence treatment strategies, as highlighted by Yoshifuji H *et al*, who showed that patients with anti-ARS-associated interstitial pneumonia respond significantly better to corticosteroids compared to “seronegative” interstitial pneumonia. However the anti-ARS-positive group were also shown to have a higher rate of relapse with recurrent lung disease (48). This suggests the need to treat patients aggressively with early intervention, with slower tapering of corticosteroids together with additional immunomodulatory therapy. From a clinical perspective, early identification of this subset of patients may influence management. Furthermore, studies described in Section 1.6.4.2 suggest a potential direct role of both candidate autoantigens and specific anti-ARS autoantibodies in disease pathogenesis that may lead to more specific targeted therapies (80, 107, 135, 164).

**Table 13: Aminoacyl-tRNA synthetases**

<b>Class I tRNA synthetases</b>	<b>Class II tRNA synthetases</b>
Arginyl	<u>Alanyl (PL-12)</u> <sup>π</sup>
Cysteinyl	<u>Asparaginyl (KS)</u> <sup>π</sup>
Glutamyl	<u>Histidyl (Jo-1)</u> <sup>π</sup>
Glutaminyl*	<u>Glycyl (EJ)</u> <sup>π</sup>
<u>Isoleucyl (OJ)*</u> <sup>π</sup>	Lysyl
Leucyl*	<u>Phenylalanyl (Zo)</u> <sup>π</sup>
Methionyl	Prolyl
Tryptophanyl	Seryl
<u>Tyrosyl (Ha)</u> <sup>π</sup>	<u>Threonyl (PL-7)</u> <sup>π</sup>
Valyl	

<sup>π</sup>autoantigenic ARS molecules

\*autoantibodies to isoleucyl (OJ) also react against other synthetases as part of a class I multi-enzyme complex including leucyl and glutaminyl-tRNA

#### **4.4.2 Anti-SAE autoantibodies**

Work described in this chapter has confirmed that SAE is an important autoantigen target in IIM where it is associated with specific clinical and associations. Anti-SAE, like anti-Mi-2, appears to be highly specific for DM. The frequency of anti-SAE in the AOMIC / RNHRD DM cohort was 8%. This is in comparison to Mi-2 which, depending on detection methods, is seen in 10-20% of DM patients (126, 128). Anti-SAE autoantibody positive DM patients have signs of hallmark cutaneous disease whereas interstitial lung involvement appears to be rare, and when present is mild. Systemic features including weight loss and dysphagia are a common manifestation of this subgroup. Furthermore, anti-SAE may be an important prognostic marker in patients who initially present with cutaneous DM. The majority of anti-SAE positive cases presented with skin lesions first and all went onto develop myositis. This finding requires further validation in larger patient groups such as those presenting with clinically-amyopathic DM under the care of dermatologists. The association with cancer and anti-SAE does not appear as strong as the now well recognised association with anti-p155/140 autoantibodies as described previously (143, 157, 158), and also in this chapter. Interestingly, the frequency anti-SAE autoantibodies with cancer-associated myositis were similar to that found for anti-Mi-2 autoantibodies in a recent study (158). In the future, larger studies will facilitate more detailed investigation of the clinical specificity of anti-SAE in comparison to other DM-specific autoantibodies.

Post-translational modifications are essential for full biological activity and allow a rapid cellular mechanism in response to changes in the environment. However, these modifications can also lead to the creation of novel self-antigens and the development of autoimmunity. Examples of this can be seen with glycosylation and hydroxylation modifications in collagen-induced arthritis, citrullination in rheumatoid arthritis and phosphorylation in SLE (165).

Small ubiquitin-like modifiers (SUMO) play a key role in the post-translational modification of specific proteins. The SUMO family consists of four members of which SUMO-1 is the best characterised. Sumoylation leads to the formation of stable conjugates of target proteins including protein kinases and transcription factors (166-169). Studies have implied a role for sumoylation in nucleo-cytoplasmic transport and signal transduction (167, 168). SUMO has been linked to certain inflammatory diseases. Franz *et al* demonstrated high levels of SUMO-1 mRNA expression in rheumatoid synovial tissue compared to controls. The relevance of

this over-expression is unclear but it was suggested that through inhibition of apoptosis, the survival of synovial fibroblasts may be increased (170). In addition, Janka *et al* have reported the presence of autoantibodies against SUMO-1 and SUMO-2 autoantigens in patients with primary biliary cirrhosis (PBC) (171). PBC is associated with other autoimmune conditions and several case reports have described PBC with co-existent myositis (172, 173).

SAE is a heterodimer composed of the subunits SAE1 and SAE2. Desterro *et al* have shown that the two polypeptides SAE1 and SAE2 have apparent molecular weights of 40 and 90 kDa respectively, with predicted molecular masses of 38 kDa and 72 kDa (174). These results are consistent with the data found in this study, where SAE2 was found to migrate at a higher molecular weight than predicted on SDS-PAGE. SAE functions in an ATP-dependant manner and forms a thioester bond between the SAE2 subunit and SUMO, prior to the transfer of SUMO to the E2 conjugating enzyme (174, 175). IIF staining using the SAE subunits have shown that SAE is distributed throughout the nuclei but excluded from the nucleoli (176). These results are consistent with the IIF staining patterns seen with anti-SAE positive sera, where a coarse speckled nucleolar sparing staining pattern was observed. The additional fine cytoplasmic speckle seen in some anti-SAE-positive cases suggests a limited cytosolic pool of the SAE, as previously shown with the sumoylation of RANGAP 1 (177).

Sumoylation is also implicated in the transcriptional repression of histone deacetylase a component of the nucleosome remodelling and deacetylase complex that also includes the myositis specific autoantigen, Mi-2 (178). Therefore, it is possible that the production of autoantibodies to either SAE or Mi-2 in DM maybe the consequence of the same post-translational modification event.

There are other protein targets that are modified by sumoylation (168). Intriguingly these include a member of the GATA family of transcription factors, GATA-2, which is expressed in adult endothelial cells. GATA-2 plays a key role in the transcriptional regulation of endothelial specific genes including endothelin-1 (ET-1), nitric oxide synthetase and Von-Willebrand factor. Both SUMO-1 and 2 covalently modify GATA-2 regulating ET-1 expression by suppressing promoter gene activity (179). This observation is of interest because endothelial cell injury and subsequent microvasculopathy in both muscle and skin lesions is well described in DM (74, 75, 180).

In summary, one can hypothesize that disruption of sumoylation of specific target proteins either via SUMO and / or the enzyme subunits SAE 1 / 2 play a role in the pathogenesis of certain DM clinical phenotypes. The clinical homogeneity of anti-SAE-positive patients, suggests that this is an important finding.

#### **4.4.3 Anti-p155/140 autoantibodies**

Over the last two years, several groups have reported an autoantibody termed anti-p155/140 based on the molecular weights of the polypeptide targets (155 and 140 kDa) in adult DM. Targoff *et al* and Kaji *et al* have identified this autoantibody specificity in 21% and 13% of DM patients respectively (143, 157). In both studies anti-p155/140 positive cases had more severe cutaneous involvement including more frequent V-sign and Shawl-sign rash, and very interestingly an association with malignancy. Furthermore, anti-p155/140 autoantibodies appear to be a negative predictor for interstitial lung disease. We have confirmed this observation in our cohort of adult DM patients. Chinoy *et al* investigated this further in larger cohort of patients from the AOMIC registry and found that the risk of malignancy in anti-p155/140 positive cases was significantly higher than anti-p155/140 negative cases (odds ratio of 23.2, 95% confidence interval 6.1-84.5) (158). These studies, and the work described in this chapter, highlight the future potential of routinely testing for anti-p155/140 to aid clinicians in the detection of cancer-associated myositis. The next step is to further validate the clinical utility of anti-p155/140, which can then be translated into routine clinical practice.

In this RNHRD cohort study only four patients out the overall IIM cohort had a history of malignancy and all were anti-p155/140 autoantibody positive. The diagnosis of cancer was either concurrent or within a few months onset of their DM, in concordance with the findings described by Kaji *et al* (157). Kaji *et al* also described two patients who were anti-p155/140-positive without malignancy and two patients who were clinically amyopathic although it is not clear if they were the same patients. Of interest, in this study, the three patients who were anti-p155/140 positive / malignancy negative also had CADM.

Preliminary work has identified the target of the anti-p155 autoantibodies as transcriptional intermediary factor 1-gamma (TIF1- $\gamma$ ) (159). Also known as Ret fused gene 7 (RFG7), this is a member of unique RING finger proteins involved in cellular differentiation (181). The discovery of the p155/140 autoantigen (TIF1- $\gamma$ ) in DM

patients with cancer-associated myositis may increase our understanding of the relationship between autoimmunity and cancer.

## **4.5 Appendix**

### **I. Acknowledgements (including work done in conjunction with others)**

Mrs J Dunphy (JO) and Mrs P Owen (PO) previously performed ANA IIF on sera from IIM cases as part of the RNHRD (Bath Institute for Rheumatic Diseases) Diagnostic Laboratory Service. I would like to thank them for giving me permission to include this data in this study.

I would also like to acknowledge and thank Dr Zoe Betteridge (ZB). The protein isolation, mass spectrometry work and immunodepletion experiments (anti-Zo and anti-SAE studies) were conducted by Dr Zoe Betteridge. I thank her for giving me permission to include her work in this thesis in conjunction with the clinical studies on anti-Zo and anti-SAE autoantibodies.

I would like to thank all the patients who have attended the RNRHD CTD clinic and given their consent to participate in these studies.



## **CHAPTER FIVE**

### **RESULTS**

#### **Serological subsets and clinical associations of myositis-specific and myositis-associated autoantibodies in juvenile dermatomyositis**

##### **5.1 Introduction**

###### **5.1.1 Epidemiology**

The juvenile idiopathic inflammatory myopathies (IIMs) are a group of rare but chronic systemic autoimmune conditions of childhood. Approximately one-fifth of all cases of IIM start in childhood, with an annual incidence of 2.5-5 cases per million population (19). Juvenile dermatomyositis (JDM) is the most common of the idiopathic inflammatory myopathies (IIM) of children, whereas juvenile polymyositis and inclusion body myositis are rarely described. The reported incidence of JDM ranges between 0.8-4.1 per million children per year (18-20). The peak age of onset is 7 years and most studies show that it is just over twice as common in girls than boys, although this ratio may be more comparable in patients under the age of 6 (19, 24). A younger age of onset may confer a worse overall prognosis and 25% cases occur before the age of 4 years (182).

###### **5.1.2 Clinical features**

JDM is chronic and potentially debilitating. Despite improvements in multi-disciplinary treatment approaches JDM is still associated with significant morbidity and mortality. Classic JDM rash is similar to adult DM lesions. Heliotrope rash, Gottron's papules over the fingers or extensor aspects of the limbs, and periungal cuticular overgrowth with capillary dilatation are prominent features. Cutaneous lesions are the presenting feature in the majority (over 50% cases), and proximal muscle weakness is the first symptoms in a quarter (183). JDM is a multi-system disease and most children will have other 'systemic' manifestations at onset, including fever, weight loss, anorexia, mouth ulcers, alopecia and irritability (24). In comparison to adult DM, vasculitis is a more prominent feature leading to various clinical signs including subcutaneous oedema and subsequent skin ulceration. Other systemic involvement includes gastrointestinal (GI) vasculitis causing abdominal pain, GI ulceration, bleeding and perforation (24, 183). Dystrophic calcinosis (in the skin, subcutaneous tissue or muscle), acquired lipodystrophy and contractures are also relatively common manifestations in JDM, and are said to be associated with chronic cutaneous inflammation, longer disease duration and delay in diagnosis / treatment onset (184, 185). JDM-scleroderma overlap (JDM-SSc) is well recognised

in JDM children with a history of Raynaud's phenomenon, sclerodactyly and other sclerodermatous skin changes. JDM can also overlap with systemic lupus erythematosus (JDM-SLE). In addition, it is well described that a large number of JDM children can also develop overlap inflammatory arthritis (usually in association with JDM-SSc or SLE) (24, 186). In contrast to adult DM, interstitial pneumonia and cancer-associated myositis are very rare in JDM (53, 55-57). Due to the heterogeneity of the condition with multi-organ involvement, clinical outcomes and prognosis are difficult to predict. Certain clinical features, such as skin ulceration, gastro-intestinal involvement (dysphagia and ischaemic ulceration from chronic endarteropathy) or respiratory disease / respiratory muscle weakness (especially leading to aspiration pneumonia) have been proposed as predictors of severe disease course in JDM (21, 25, 57, 187).

### **5.1.2 Clinical Assessment**

There are now standardised validated clinical tools used in the assessment of JDM both in clinical practice and in clinical trials. These include the Childhood Myositis Assessment Scale (CMAS) and the Childhood Health Assessment Questionnaire (CHAQ) (188-191). Additional measures of disease activity include the Physicians' global assessment using a visual analogue scale (VAS) or Likert scale, and a patient / parent global assessment of overall well-being using the same method (VAS) (192). In addition, the Cutaneous Assessment tool (CAT) has been partially validated for the assessment of skin disease in JDM (193, 194).

Standard laboratory markers include muscle enzymes; creatinine kinase (CK), lactate dehydrogenase (LDH), aldolase, and other inflammatory markers including C-reactive protein (CRP) or erythrocyte sedimentation rate (ESR). Magnetic resonance imaging (MRI) of proximal muscle groups is now being frequently used for diagnostic purposes as well assessment of muscle disease activity, and in some cases may be preferred to muscle biopsy. T2 fat suppressed images can demonstrate muscle, subcutaneous and cutaneous inflammation and oedema. T1 weighted images are used to establish the degree of fatty infiltration and muscle atrophy, which is representative of damaged muscle (195-197).

***Childhood Myositis Assessment Scale (CMAS) (188, 189)***

Assessment comprises of a series of timed tasks testing muscle strength, endurance and function. See Chapter 5.5: Appendix I.

***Childhood Health Assessment Questionnaire (CHAQ) (190, 191)***

The CHAQ is a functional questionnaire completed by the patient and / or parent to assess activities of daily living. See Chapter 5.5: Appendix I.

***Cutaneous Assessment tool (CAT) (193, 194)***

The CAT is used in both adult and juvenile DM to assess skin disease measuring both activity and damage, using a *priori* weighting scale. See Chapter 5.5: Appendix I.

**5.1.3 Aetiology and pathogenesis**

The aetiology of JDM remains unclear, however similar to adult IIM, it is well recognised that genetic susceptibility combined with environmental triggers leads to the development of autoimmunity and thus immune dysfunction. The strongest genetic association for autoimmune connective tissue diseases, including JDM is the human leucocyte region (HLA) within the major histocompatibility complex (MHC). The MHC region consists of clustered multiple alleles coding for proteins central to the immune system. The 8.1 ancestral MHC haplotype, (HLA-B\*08-DRB1\*03-DQA1\*05-DQB1\*02) is associated with JDM, where the strongest signal appears to come from the HLA class I region (10, 198-200). Recent work demonstrates that the DQA1\*0301 allele confers an additional risk for JDM, whereas certain alleles, DQA1\*0201, DQA1\*0101 and DQA1\*0102 appear protective (200). Certain cytokine polymorphisms, including  $\text{TNF}\alpha$ -0308A and IL-1 receptor antagonist, are also associated with Caucasian JDM with a higher risk of cutaneous ulceration and calcinosis (201, 202).

Environmental factors appear to play a role in the development of JDM. Reports suggest a temporal association to the disease onset. Various infectious agents have been implicated with the onset of JDM, in particular Group A streptococcus and several viruses including coxsackie, enterovirus and parvovirus (203-206). Seasonal birth patterns have been assessed in juvenile-onset IIM, in comparison to patients with adult-onset DM and healthy controls. It was noted that juvenile-onset IIM Hispanic patients had a seasonal birth pattern significantly different to controls. Birth patterns also differed in patients with certain HLA risk factors including

DRB1\*0301 and DQA1\*0501 alleles. The authors concluded that birth distributions appear to have a stronger seasonality in juvenile IIM, suggesting a role for perinatal or early-life exposures (95).

Similar to adult IIM, both cellular and humoral mechanisms are involved in pathogenesis of JDM, and recent evidence suggests a role of the innate immune system (17, 73, 74, 207-210). Antibody-mediated complement deposition in the vessels plus dendritic cell (DCs), CD4+ T cell and B cell perivascular and perifascicular infiltration and upregulation of MHC class I and class II molecules by myocytes are characteristic. Vasculopathy plays a major pathophysiological role in JDM. Affected muscle is characterised by endothelial dysfunction and capillary dropout. Clinical features in the skin and GI tract suggest similar pathology (73). Hallmark features on muscle biopsy include oedema of the capillary endothelial wall with subsequent occlusive vasculopathy, perivascular and perimysial inflammatory cell infiltrate, muscle fibre changes including perifascicular atrophy. Evidence of muscle degeneration and regeneration is present with neonatal myosin. Recently an international consensus scoring system for the analysis of JDM muscle biopsy specimens has been published (211). Recent work has highlighted the importance of type 1 interferons (IFN) in JDM disease mechanisms. Type I IFN is believed to be critical in the propagation of autoimmunity, and this mechanism has been implicated in the pathogenesis of JDM as well as adult DM (212, 213). Studies using microarray technology have demonstrated up-regulation of type-1 IFN induced genes in DM (214, 215). DM muscle contains high numbers of gene transcripts associated with type 1 IFN, the source of which are CD4+ plasmacytoid dendritic cells (pDCs) that have chemokine activity. Similar findings were also noted in DM skin. Finally, the potential role of maternal microchimerism in JDM has been highlighted over the last few years. The presence of maternal cells in peripheral blood and diseased muscle of JDM cases may induce autoimmunity i.e. 'a graft-versus-host disease' mechanism (216-218).

#### **5.1.4 Autoantibodies in JDM and JDM-overlap syndromes**

Until recently, unlike adult IIM, specific autoantibodies in juvenile IIM especially JDM, were detected infrequently despite the majority being ANA positive on IIF. As in adult IIM, autoantibodies are divided into MSAs and MAAs, the latter typically described in JDM-overlap syndromes (JDM-SSc and JDM-SLE).

Unlike adults the frequency of anti-aminoacyl tRNA synthetase (ARS) autoantibodies in juvenile myositis, in particular JDM, is much lower. Rider *et al* used immunodiffusion and immunoprecipitation (IPP) to screen sera from 77 children with myositis and overlap CTD, and anti-ARS were detected in only 2.6% cases (125). Using the same techniques, Feldman *et al* investigated 42 children; thirty five with JDM and 7 with other forms of IIM, and none had anti-ARS autoantibodies (127). The clinical features of anti-ARS-positive JDM cases are similar to their adult counterparts: moderate to severe weakness, arthritis, Raynaud's phenomenon, mechanic's hands, fevers and interstitial lung disease.

Similar to the ARS molecules, SRP is rarely targeted by autoantibodies in juvenile myositis. Based on previous studies the frequency of anti-SRP is 1% (125, 127). Recently, Rouster-Stevens *et al* described three cases of juvenile anti-SRP myositis in their cohort of 123 children and based on their clinical features the patients were comparable to adults with the same autoantibody specificity. In particular; all three children had severe weakness, very high creatinine kinase (CK) levels, were refractory to treatment and had myofibre necrosis on biopsy (219). Of interest, the ethnic origin of these patients was African American and Love *et al* has previously reported a higher frequency of anti-SRP in adult African American cases (6).

The reported frequency of anti-Mi-2 autoantibodies in JDM or JDM overlap is up to 5% (125, 127). Like adult DM, this autoantibody specificity is described in children with hallmark cutaneous DM lesions including Gottron's papules, heliotrope rash, cuticular overgrowth and rashes on the neck and trunk. Moreover, patients may have milder muscle involvement, appear to have a monocyclic clinical course and respond well to therapy (125-127).

MAAs: anti-U1-RNP, anti-PM-Scl, anti-U3-RNP, anti-Ku and anti-topoisomerase are typically detected in children with JDM associated with systemic sclerosis features including scleroderma skin changes and Raynaud's. Based on previous data, collectively the overall frequency in JDM / JDM-overlap is between 15-20% (57, 220, 221).

Previously, the overall frequency of defined MSAs and MAAs in children with IIM was between 20-40%, however using full serological testing, specifically IPP this frequency is now much higher following the studies described in the next two

chapters of this thesis, and earlier reports from Wedderburn *et al*, Targoff *et al* and Oddis *et al* (10, 143, 222).

The work described in this chapter is a clinical and serological study of JDM and JDM-overlap children recruited to the JDRR (UK and Ireland). Chapter 6 describes the work on novel autoantibody specificities in JDM.

## **5.2 Patients and Methods**

### **5.2.1 Patients and sera**

Subjects for the study described in this chapter were recruited from:

- The Juvenile Dermatomyositis Registry and Repository (JDRR), UK and Ireland (24).

The JDRR has recruited patients from 10 centres around the UK (for details see Chapter 5, Appendix III) with juvenile-onset myositis, below the age of 16 years at disease onset and diagnosis. All JDM cases had probable or definite disease according to *Bohan and Peter* criteria (1, 2). JDM-scleroderma overlap (JDM-SSc) was defined as JDM children with a history of Raynaud's phenomenon, sclerodactyly and other sclerodermatous skin changes (two or more of the above features). Using a standardised proforma demographic and clinical data were recorded at diagnosis and prospectively at subsequent visits, on average every 6 months (see Chapter 2, Section 2.1.2 and 2.1.3, and appendix). The clinical information recorded included specific cutaneous manifestations including the presence of Gottron's lesions, skin ulceration, oedema, calcinosis and the distribution of skin rash over the body. Details on muscle involvement included muscle enzymes (CK and LDH) and CMAS at disease baseline. The development of malignancy (within three years of diagnosis) was not specifically recorded on the clinical proforma. However, clinicians were able to enter free text data on clinically significant events during disease course, which was therefore likely to capture this information. Data were stored using anonymous codes onto a central database. Serum samples were taken at the time of diagnosis and stored at -20°C until required.

### **5.2.2 Serological Methods**

#### ***ANA-Immunofluorescence (IIF)***

IIF was previously performed by standard methods using HEp-2 cells and fluorescein-labelled anti-human IgG immunoglobulin (Sigma, UK).

### ***Protein Radio-immunoprecipitation (IPP)***

Sera stored at -20°C were thawed at room temperature. IPP from K562 cell extracts was performed as described in Chapter 2, see Sections 2.3.2 and 2.3.3. Briefly, 10 µl sera was mixed with 2 mg protein-A-Sepharose beads (Sigma, UK) in IPP buffer (10 mM Tris-Cl pH 8.0, 500 mM NaCl, 0.1% v/v Igepal) at room temperature for 30 min. Beads were washed in IPP buffer prior to the addition of 120 µl [<sup>35</sup>S] methionine labelled K562 cell extract. Samples were mixed at 4°C for 2 hr. Beads were washed in IPP buffer followed by TBS buffer (10 mM Tris-Cl pH 7.4, 150 mM NaCl) before being resuspended in 50 µl SDS sample buffer (Sigma, UK). After heating, proteins were fractionated by 10% SDS-PAGE, enhanced, fixed and dried. Labelled proteins were analysed by autoradiography.

### **5.2.3 Ethical approval**

All patients gave fully informed written / parental consent to participate and provide biological samples according to the Declaration of Helsinki under both national multi-centre and local ethical committee regulations.

### **5.2.4 Statistical Analysis**

Clinical associations were derived from 2×2 contingency tables using the chi-squared test, or two-tailed Fisher's exact test where individual cells valued five or less. *P* values less than 0.05 were considered as significant (uncorrected *P* values presented in view of small numbers in each autoantibody group). SPSS for Windows (version 14) was used to perform statistical analysis.

## **5.3 Results**

Serum samples for serological typing were available from 162 children recruited to the registry. Full clinical data were available on 160 children and 74% were female. The median age of disease onset was 6 years, inter-quartile range (IQR) 3-9 and median age at diagnosis was 7 years (IQR 4, 10). One hundred and twenty seven cases were Caucasian and 35 cases were non-Caucasian.

The median follow-up from disease onset to the time of data analysis for this study was 48 months (IQR 33, 72) for the overall cohort.

One hundred and thirty seven children had JDM. In this study, 21 children were defined as JDM-SSc. Two children were defined as having indeterminate juvenile myositis, not specifically JDM or JDM-SSc.

### **5.3.1 Serological profiles**

Following ANA IIF and IPP, autoantibody specificity was categorised into recognised MSAs, myositis-associated autoantibodies (MAAs), unidentified / novel or negative. The overall frequency of previously defined MSAs and MAAs in this cohort study is shown in Table 14. Table 15 shows a breakdown of serological results and ethnicity (Caucasian and non-Caucasian).

#### ***IPP results***

Thirteen out of 162 (8%) cases were positive for previously described MSAs: anti-Jo-1 (1.2%) and anti-Mi-2 (6.8%). No cases were positive for anti-SRP autoantibodies. Anti-Mi-2 was found exclusively in JDM patients, with a frequency of 7.9%, and not in any JDM-overlap patients. Anti-Jo-1 was found in JDM-overlap only. The frequency of 'known' MSA-positive in JDM was 8% compared to 9.5% in JDM-overlap.

The overall frequency of MAAs in this study cohort was 14.2%: anti-U1-RNP (5.6%), anti-PM-Scl (6.8%), anti-U3-RNP (0.6%), anti-Ku (0.6%) and anti-Topoisomerase (0.6%). There was a much higher frequency of MAA-positive cases in JDM-overlap (81%) compared to MAA-positive JDM (4.4%). Of note, anti-U1-RNP and anti-PM-Scl was found in higher frequency in Caucasians compared to non-Caucasian children.

Novel autoantibodies were detected in 40% of the overall cohort, and these specificities were found exclusively in JDM cases with a frequency of 47.4% (this work is described in detail in Chapter 6). In the overall cohort, 37.6% of cases were either negative or had undefined bands on IPP.

See Figures 15A-C for examples of MAAs and MSAs in JDM study.

For full IPP results see Chapter 5.5, Appendix II.

#### ***ANA IIF results***

For overall results see Chapter 5.5, Appendix II.

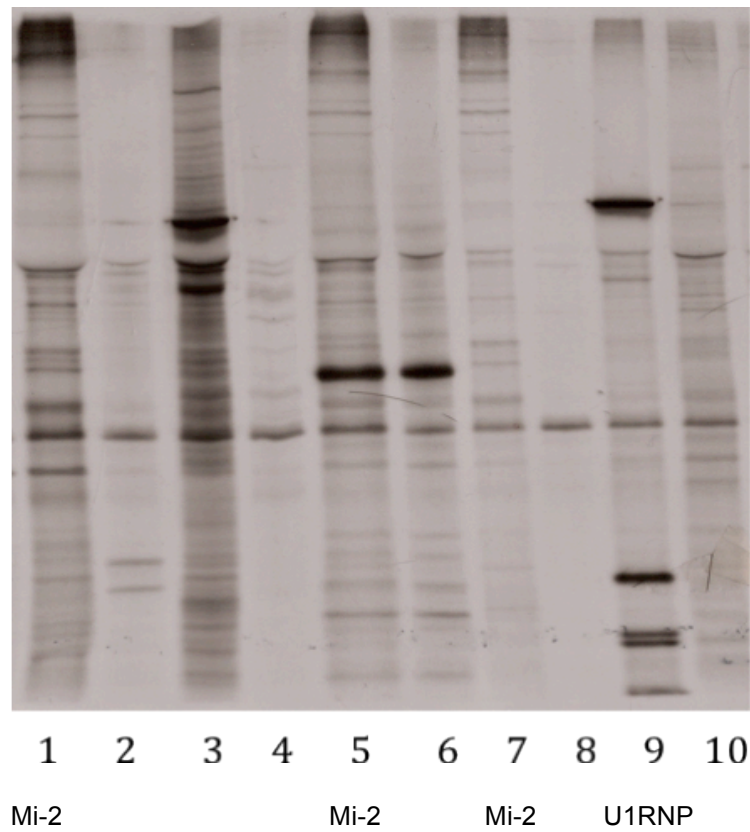
In total, ANA IIF was performed on 124 samples (there were insufficient sera in 38 samples for both ANA IIF and IPP, therefore only IPP was performed). Out of 124 cases, 96 (77.4%) were ANA positive by IIF.



**Figure 15A: Immunoprecipitation of MSAs and MAAs in JDM cohort study**

10% SDS PAGE of immunoprecipitates of [ $^{35}$ S] labelled K562 cell extracts.

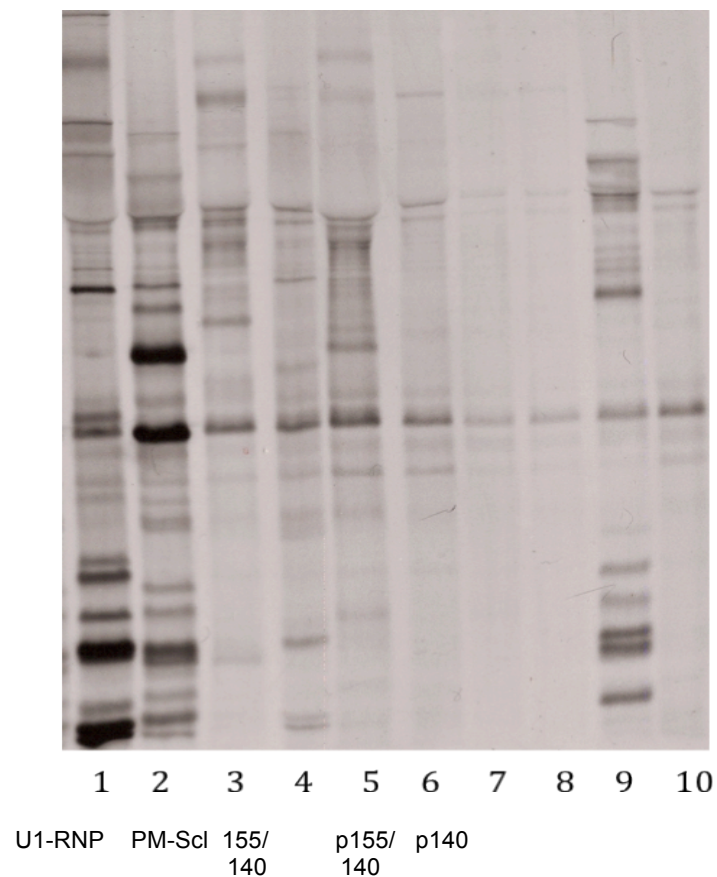
JDM sera used for immunoprecipitation: Lane 1; anti-Mi-2, Lane 5; anti-Mi-2, Lane 7; anti-Mi-2, Lane 9; anti-U1-RNP.



**Figure 15B: Immunoprecipitation of MSAs and MAAs in JDM cohort study**

10% SDS PAGE of immunoprecipitates of [ $^{35}$ S] labelled K562 cell extracts.

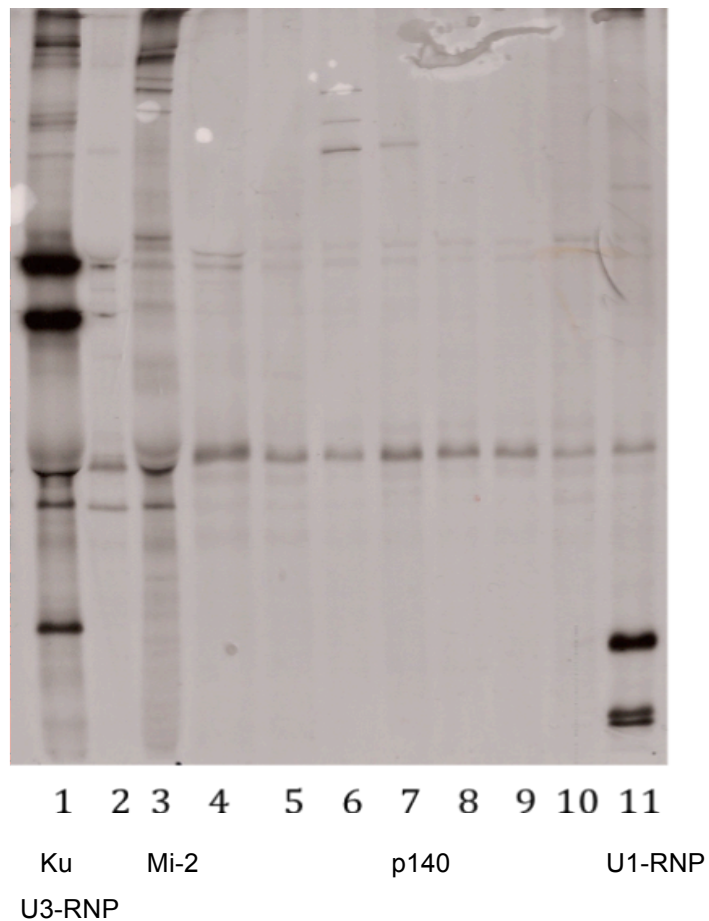
JDM sera used for immunoprecipitation: Lane 1; anti-U1-RNP, Lanes 2 and 9; anti-PM-Scl, Lane 3 and 5; novel anti-p155/140 (see Chapter 6), Lane 4; possible anti-U1-RNP, Lane 6; novel anti-p140 (see Chapter 6).



**Figure 15C: Immunoprecipitation of MSAs and MAAs in JDM cohort study**

10% SDS PAGE of immunoprecipitates of [<sup>35</sup>S] labelled K562 cell extracts.

JDM sera used for immunoprecipitation: Lane 1; anti-Ku and anti-U3-RNP, Lane 3; anti-Mi-2, Lane 7; novel anti-p140 (see Chapter 6), Lane 11; anti-U1-RNP.



**Table 14: Myositis-specific autoantibodies and Myositis-associated autoantibodies in JDM and JDM-overlap cohort**

<b>Autoantibody</b>	<b>JDM n=137</b>	<b>JDM-overlap n=21</b>	<b>Other myositis n=2</b>	<b>Overall n=162</b>
<b>MSAs</b>				
Anti-ARS				
Anti-Jo-1	0	9.5	0	1.2*
Anti-PL-12	0	0	0	0
Anti-PL-7	0	0	0	0
Anti-EJ	0	0	0	0
Anti-OJ	0	0	0	0
Anti-KS	0	0	0	0
Anti-Ha	0	0	0	0
Anti-Zo	0	0	0	0
Anti-SRP	0	0	0	0
Anti-Mi-2	7.9	0	0	6.8
<b>MAAs</b>				
Anti-U1-RNP	2.2	28.6	0	5.6
Anti-PM-Scl	2.2	38.1	0	6.8
Anti-U3-RNP	0	4.8	0	0.6
Anti-Ku	0	4.8	0	0.6
Anti-Topo	0	4.8	0	0.6
<b>Novel antibodies</b>	47.4	0	0	40.1
See Chapter 6				
<b>Unknown bands</b>	16.1	9.5	50	15.4
<b>Negative</b>	25.5	0	50	22.2

Values shown are in percentages

MSAs: myositis-specific autoantibodies

MAAs: myositis-associated autoantibodies

Topo: Topoisomerase (Scl-70)

\*One anti-Jo-1-positive case also positive for anti-Ro autoantibodies

**Table 15: Myositis-specific autoantibodies and Myositis-associated autoantibodies in JDM and JDM-overlap cohort (ethnic groups)**

<b>Autoantibody</b>	<b>JDM Caucasian n=108</b>	<b>JDM Non-Caucasian n=31</b>	<b>JDM-overlap Caucasian n=19</b>	<b>JDM- overlap Non- Caucasian n=4</b>
<b>MSAs</b>				
Anti-ARS (Jo-1)	0	0	10.5	0
Anti-SRP	0	0	0	0
Anti-Mi-2	8	6.5	0	0
<b>MAAs</b>				
Anti-U1-RNP	1.9	0	31.6	25
Anti-PM-Scl	0.8	6.5	42.1	0
Anti-U3-RNP	0	0	5.3	0
Anti-Ku	0	0	0	25
Anti-Topo	0	0	5.3	0
<b>Novel antibodies</b> See Chapter 6	42.6	61.3	0	0
<b>Unknown bands</b>	17.6	9.7	5.3	25
<b>Negative</b>	27.8	16.1	0	0

Values shown are in percentages

MSAs: myositis-specific autoantibodies / MAAs: myositis-associated autoantibodies

Topo: Topoisomerase (Scl-70)

### 5.3.2 Clinical associations of myositis autoantibodies in study cohort

The overall clinical features of the entire cohort are shown in Table 16. The main clinical features of MSAs (anti-Jo-1 and –Mi-2) and MAAs are shown in Table 17. Two cases were anti-Jo-1-positive with JDM-overlap. The median age at onset was 9 years and both cases were females. Both children had DM rash with myositis, arthritis, sclerodermatous puffy fingers and Raynaud's phenomenon. This MSA subset had no complicating features such as skin ulceration or calcinosis, however the numbers are small. One anti-Jo-1-positive child also had interstitial pneumonia and mechanic's hands.

Anti-Mi-2 was detected in 11 children and all had JDM with hallmark cutaneous changes and a wide distribution of rash: Gottron's papules (100%), cutaneous oedema 63%, periorbital (100%) and periungal overgrowth with capillary changes (91%). The median age at onset was 12.5 years and 90% cases were girls.

MAAs (specifically anti-U1-RNP, -PM-Scl, -U3-RNP, -Ku, -Topo) were detected in 17 children with JDM-overlap and 6 children with a diagnosis of JDM. MAA-positive children had a median age at onset of 9 years, and 78% were female. Sixty-five% of MAA-positive cases had Gottron's papules, with periorbital rash (35%), periungal changes (57%) and rash on the trunk (17%). Other features were frequent: arthritis (65%), scleroderma skin changes (61%), Raynaud's (48%), calcinosis (21%) and lipoatrophy (35%). Both anti-PM-Scl and anti-U1-RNP autoantibodies were present in 5-7% of juvenile IIM respectively (a similar frequency to anti-Mi-2).

When comparing anti-Mi-2-positive children with MAA-positive children the frequency of skin lesions - Gottron's papules ( $P=0.034$ ) and cutaneous oedema ( $P=0.021$ ), distribution of rash – periorbital ( $P<0.001$ ), were significantly higher in the anti-Mi-2 group. Sclerodermatous features were significantly more frequent in the MAA-positive group ( $P<0.001$ ).

**Table 16: Overall clinical features of the UK JDM Cohort study (n=160)**

<b>Clinical feature</b>	<b>Frequency (%)</b>
Female	73.8
Caucasian	79.4
<b>Skin disease</b>	
Gottron's	82.9
Ulceration	25.3
Lipoatrophy	14
Oedema	34.8
Calcinosis	23.3
<b>Distribution of rash</b>	
Periorbital	71.4
Periungal	66.2
Trunk	14.3
Small joints	70.8
Large joints	51.3
<b>Other features</b>	
Arthritis	44.7
Sclerodermatous features	12.6
Raynaud's phenomenon	16.6
Dysphagia	31.4
Mouth ulcers	27
Alopecia	29.7
Interstitial pneumonia	5

Values are in percentages

**Table 17: Clinical features of MSAs and MAAs in JDM and JDM-overlap cohort**

	Anti-Jo-1 n=2 (%)	Anti-Mi-2 n=11 (%)	MAA n=23 (%)
<b>Median age (onset)</b>	9	12.5	7
<b>Median age (diagnosis)</b>	9	12.5	9
<b>Female</b>	100	90	78
<b>Type of skin lesion</b>			
Gottron's papules	100	100 *	65 *
Ulceration	0	9	13
Oedema	0	63 **	17 **
Calcinosis	0	9	21
Lipoatrophy	0	9	35
Mechanic's hands	50	0	0
<b>Rash distribution</b>			
Periorbital	100	100 δ	35 δ
Periungal	100	91	57
Small joints	100	82	65
Large joints	50	45	30
Trunk	0	0	17
<b>Other features</b>			
Arthritis	100	36	65
Scleroderma	100	0 α	61α
Raynaud's	100	9	48
Dysphagia	0	45	30
Mouth ulcers	0	36 αα	4 αα
Alopecia	100	30	26
Interstitial pneumonia	50	0	4
<b>Myositis</b>			
Raised CK	100	70	76

Values are in percentages, unless otherwise stated, (n/a: full data not available)

\*  $P_{(uncorr)}=0.034$ : anti-Mi-2-positive *versus* MAA-positive

\*\*  $P_{(uncorr)}=0.021$ : anti-Mi-2-positive *versus* MAA-positive

δ  $P<0.001$ : anti-Mi-2-positive *versus* MAA -positive

α  $P<0.001$ : anti-Mi-2-positive *versus* MAA-positive

αα  $P_{(uncorr)}=0.05$ : anti-Mi-2-positive *versus* MAA-positive



## 5.4 Discussion

This chapter describes a serological study of JDM and JDM-overlap in children recruited to the JDERR. The aim of the work in this Chapter was to establish the frequency of known MSAs and MAAs, and describe the clinical associations in this JDM UK cohort.

In comparison to previous serological studies in juvenile IIM summarised in several review articles (221, 223, 224), the overall frequency of MSAs and MAAs are similar, especially the anti-Mi-2 and MAAs subsets. In a recent study by Espada *et al*, 64 Argentine Caucasian children with juvenile IIM (including 40 JDM cases and 17 myositis overlap cases) were screened using IPP and RNA immunoprecipitation. The frequency of anti-Mi-2 was 6%, anti-PM-Scl was 3% and anti-U1-RNP autoantibody was 11%. No children were positive for anti-ARS autoantibodies (225). In this cohort study, the frequency of anti-ARS autoantibodies was generally less than previously described (125, 127), with less anti-Jo-1-positives and no children testing positive for non-Jo-1-ARS. In contrast to adult IIM where autoantibodies to ARS are a common target (as described in Chapter 3), this group of MSAs are detected infrequently in juvenile myositis. Although the numbers of children with anti-ARS autoantibodies were small, the clinical manifestations do resemble that of their adult counterparts with the anti-synthetase syndrome (see Chapter 3) (46). Anti-SRP autoantibodies were not detected in this cohort, which may reflect the ethnic make-up of this cohort (the majority of cases being Caucasian) with no cases of 'juvenile PM' being recruited to the JDERR at the time of this study. Typically anti-SRP-positivity is rare in juvenile IIM anyway and a recent report has highlighted how this MSA may be generally more prevalent in Afro-American children (219). Anti-Mi-2 was detected in JDM children with hallmark cutaneous changes, a wide distribution of rash, and a relatively higher frequency of cutaneous / subcutaneous oedema in comparison to the anti-Jo-1 and MAA subgroups. In contrast, extra-muscular / cutaneous complications were relatively less frequent except for dysphagia. The frequency of anti-Mi-2 in both juvenile and adult myositis is similar.

The frequency of MAAs in this juvenile cohort is similar to that seen in adult IIM, especially when comparing JDM-SSc patients with adult myositis/CTD overlap (139). The most common MAA specificities were anti-U1-RNP and anti-PM-Scl, with small number of children testing positive for anti-U3-RNP, -Topo, and -Ku. There were no patients with other systemic sclerosis-associated autoantibodies e.g. anti-centromere or anti-RNA polymerase I, II or III, which highlights differences between

juvenile and adult myositis-SSc overlap populations (139, 226). Children with SSc, including the diffuse form of the disease, have similar serological profiles. Anti-PM-Scl and anti-U1-RNP autoantibodies are detected in a higher frequency compared to adult onset SSc. Other specificities including anti-centromere, anti-RNA polymerase and anti-topoisomerase are more common in adult SSc (226, 227). In general, MAA-positive children had less classic skin DM lesions but myositis with more arthritis, scleroderma skin changes, Raynaud's and lipoatrophy. In comparison to anti-Jo-1 and anti-Mi-2-positive children, MAA-positives had a relatively higher frequency of calcinosis, and this observation may reflect the predominant overlap features with scleroderma. In general, overlap features are seen in about approximately 10% of juvenile-onset IIM (21, 22). In this study, 12% of children were defined as having JDM-SSc overlap, and although full data was not available to make any conclusions regarding severity of muscle disease, it has been observed that children with overlap may have milder muscle disease with lower disease activity markers (21).

In summary, the overall frequency of recognised MSA or MAAs in this cohort was around 20%. Therefore, most children with myositis are either autoantibody negative or have autoantibodies that target novel autoantigens. The majority have novel specificities and this work is described in Chapter 6.

## 5.5 Appendix I

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### CHILDHOOD MYOSITIS ASSESSMENT SCALE (CMAS) SCORING SHEET

1. **HEAD LIFT:**  
0 = Unable      3 = 30-59  
1 = 1-9 sec    4 = 60-119 sec  
2 = 10-29      5 =  $\geq 2$  min      # of sec \_\_\_\_\_
2. **LEG RAISE/TOUCH OBJECT:**  
0 = Unable to lift leg off table.  
1 = Able to clear table, but cannot touch object (examiner's hand).  
2 = Able to lift leg high enough to touch object (examiner's hand).
3. **STRAIGHT LEG LIFT/DURATION:**  
0 = Unable      3 = 30-59 sec  
1 = 1-9 sec    4 = 60-119 sec  
2 = 10-29 sec    5 =  $\geq 2$  min      # of sec \_\_\_\_\_
4. **SUPINE TO PRONE:**  
0 = Unable. Has difficulty even turning onto side; able to pull right arm under torso only slightly or not at all.  
1 = Turns onto side fairly easily, but cannot fully free right arm and is unable to fully assume a prone position.  
2 = Easily turns onto side; has some difficulty freeing arm, but fully frees arm and fully assumes a prone position.  
3 = Easily turns over, fully frees right arm with no difficulty.
5. **SIT-UPS:**  
Hands on thighs, with counterbalance \_\_\_\_\_  
Hands across chest, with counterbalance \_\_\_\_\_  
Hands behind head, with counterbalance \_\_\_\_\_  
Hands on thighs, without counterbalance \_\_\_\_\_  
Hands across chest, without counterbalance \_\_\_\_\_  
Hands behind head, without counterbalance \_\_\_\_\_  
Total Sit-up Score (0-6) \_\_\_\_\_
6. **SUPINE TO SIT:**  
0 = Unable by self.  
1 = Much difficulty. Very slow, struggles greatly, barely makes it. Almost unable.  
2 = Some difficulty. Able, but is somewhat slow, struggles some.  
3 = No difficulty.
7. **ARM RAISE/STRAIGHTEN:**  
0 = Cannot raise wrists up to the level of the A-C joint.  
1 = Can raise wrists at least up to the level of the A-C joint, but not above top of head.  
2 = Can raise wrists above top of head, but cannot raise arms straight above head so that elbows are in full extension.  
3 = Can raise arms straight above head so that elbows are in full extension.
8. **ARM RAISE/DURATION:** Can maintain wrists above top of head for:  
0 = Unable      3 = 30-59 sec  
1 = 1-9 sec    4 =  $\geq 60$  sec  
2 = 10-29 sec      # of sec \_\_\_\_\_
9. **FLOOR SIT:** Going from a standing position to a sitting position on the floor:  
0 = Unable. Afraid to even try, even if allowed to use a chair for support. Child fears that he/she will collapse, fall into a sit, or harm self.  
1 = Much difficulty. Able, but needs to hold onto a chair for support during descent. Unable, or unwilling to try if not allowed to use a chair for support.  
2 = Some difficulty. Can go from stand to sit without using a chair for support, but has at least some difficulty during descent. May need Gower's. Descends somewhat slowly and/or apprehensively; may not have full control or balance as maneuvers into a sit.  
3 = No difficulty. Requires no compensatory maneuvering.
10. **ALL FOURS MANEUVER:**  
0 = Unable to go from a prone to an all-fours position.  
1 = Barely able to assume and maintain an all-fours position. Unable to raise head to look straight ahead.  
2 = Can maintain all-fours position with back straight and head raised (so as to look straight ahead). But, cannot creep (crawl) forward.  
3 = Can maintain all-fours, look straight ahead and creep (crawl) forward.  
4 = Maintains balance while lifting and extending one leg.
11. **FLOOR RISE:** Going from a kneeling position on the floor to a standing position:  
0 = Unable, even if allowed to use a chair for support.  
1 = Much difficulty. Able, but needs to use a chair for support. (Unable if not allowed to use a chair.)  
2 = Moderate difficulty. Able to get up without using a chair for support, but needs to place one or both hands on thighs/knees or floor. (Unable without using hands.)  
3 = Mild difficulty. Does not need to place hands on knees, thighs or floor, but has at least some difficulty during ascent.  
4 = No difficulty.
12. **CHAIR RISE:**  
0 = Unable to rise up from chair, even if allowed to place hands on sides of chair seat.  
1 = Much difficulty. Able, but needs to place hands on sides of seat. Unable if not allowed to place hands on sides of seat.  
2 = Moderate difficulty. Able, but needs to place hands on knees/thighs. Does not need to place hands on sides of seat.  
3 = Mild difficulty. Does not need to place hands on seat, knees or thighs but has at least some difficulty during ascent.  
4 = No difficulty.
13. **STOOL STEP:**  
0 = Unable.  
1 = Much difficulty. Able, but needs to place one hand on exam table (or examiner's hand).  
2 = Some difficulty. Able, does not need to use exam table for support, but needs to use hand on knee/thigh.  
3 = Able. Does not need to use exam table or hand on knee/thigh.
14. **PICK-UP:**  
0 = Unable to bend over and pick up pencil off floor.  
1 = Much difficulty. Able, but relies heavily on support gained by placing hands on knees/thighs.  
2 = Some difficulty. Has some difficulty (but not "much-difficulty"). Needs to at least minimally and briefly place hand(s) on knees/thighs for support. Is somewhat slow.  
3 = No difficulty. No compensatory maneuver necessary.

The maximum possible total score for the 14 maneuvers is 52 (52 "points of muscle strength/function").

PATIENT \_\_\_\_\_ DATE \_\_\_\_\_

TOTAL CMAS SCORE: \_\_\_\_\_

## 5.5 Appendix I

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(<http://www.rheumatology.org/sections/pediatric>)

CHILDHOOD HEALTH ASSESSMENT QUESTIONNAIRE						
1						
2	<p>In this section we are interested in learning how your child's illness affects his/her ability to function in daily life. Please feel free to add any comments on the back of this page. In the following questions, please check the one response which best describes your child's usual activities (averaged over an entire day) <u>OVER THE PAST WEEK</u>. ONLY NOTE THOSE DIFFICULTIES OR LIMITATIONS WHICH ARE DUE TO ILLNESS. If most children at your child's age are not expected to do a certain activity, please mark it as "Not Applicable". For example, if your child has difficulty in doing a certain activity or is unable to do it because he/she is too young but not because he/she is RESTRICTED BY ILLNESS, please mark it as "NOT Applicable".</p>					
3		Without ANY Difficulty	With SOME Difficulty	With MUCH Difficulty	UNABLE To do	Not Applicable
4	<b>DRESSING &amp; GROOMING</b>					
5	Is your child able to:					
6	- Dress, including tying shoelaces and doing buttons?	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
7	- Shampoo his/her hair?	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
8	- Remove socks?	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
9	- Cut fingernails?	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
10	<b>ARISING</b>					
11	Is your child able to:					
12	- Stand up from a low chair or floor?	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
13	- Get in and out of bed or stand up in a crib?	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
14	<b>EATING</b>					
15	Is your child able to:					
16	- Cut his/her own meat?	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
17	- Lift up a cup or glass to mouth?	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
18	- Open a new cereal box?	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
19	<b>WALKING</b>					
20	Is your child able to:					
21	- Walk outdoors on flat ground?	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
22	- Climb up five steps?	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
23	* Please check any AIDS or DEVICES that your child usually uses for any of the above activities:					
24	- Cane	<input type="checkbox"/>	- Devices used for dressing (button hook, zipper pull, long-handled shoe horn, etc.)			<input type="checkbox"/>
25	- Walker	<input type="checkbox"/>	- Built up pencil or special utensils			<input type="checkbox"/>
26	- Crutches	<input type="checkbox"/>	- Special or built up chair			<input type="checkbox"/>
27	- Wheelchair	<input type="checkbox"/>	- Other (Specify: _____)			<input type="checkbox"/>
28	* Please check any categories for which your child usually needs help from another person BECAUSE OF ILLNESS:					
29	- Dressing and Grooming	<input type="checkbox"/>	- Eating			<input type="checkbox"/>
30	- Arising	<input type="checkbox"/>	- Walking			<input type="checkbox"/>

	Without ANY Difficulty	With SOME Difficulty	With MUCH Difficulty	UNABLE To do	Not Applicable
31					
32	<b>HYGIENE</b>				
33	Is your child able to:				
34	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
35	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
36	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
37	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
38	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
39	<b>REACH</b>				
40	Is your child able to:				
41	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
42	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
43	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
44	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
45	<b>GRIP</b>				
46	Is your child able to:				
47	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
48	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
49	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
50	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
51	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
52	<b>ACTIVITIES</b>				
53	Is your child able to:				
54	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
55	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
56	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
57	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
58	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
59	* Please check any AIDS or DEVICES that your child usually uses for any of the above activities:				
60	<input type="checkbox"/>	- Bath tub bar			<input type="checkbox"/>
61	<input type="checkbox"/>	- Long-handled appliances for reach			<input type="checkbox"/>
62	<input type="checkbox"/>	- Long-handled appliances in bathroom			<input type="checkbox"/>
63	* Please check any categories for which your child usually needs help from another person BECAUSE OF ILLNESS:				
64	<input type="checkbox"/>	- Gripping and opening things			<input type="checkbox"/>
65	<input type="checkbox"/>	- Errands and chores			<input type="checkbox"/>
66	PAIN: We are also interested in learning whether or not your child has been affected by pain because of his or her illness. How much pain do you think your child has had because of his/her illness IN THE PAST WEEK? Place a mark on the line below, to indicate the severity of the pain				
67	No pain 0  -----  100 Very severe pain				
68	GLOBAL EVALUATION: Considering all the ways that arthritis affects your child, rate how he/she is doing by placing a single mark on the line below.				
69	Very well 0  -----  100 Very poor				

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## **5.5 Appendix I**

### **Cutaneous Assessment Tool**

**Full proforma available on:**

**<http://dir-apps.niehs.nih.gov/imacs/index.cfm?action1/4home.main>**

## 5.5 Appendix II – Subjects, ANA IIF and IPP results

Study ID	Diagnosis	ANA IIF result	IPP result
UK01PB0001	JDM	Negative	140 kDa band *
UK01SS0002	JDM	Negative	Negative
UK01JR0003	JDM	Weak Homogeneous	155/140 kDa bands *
UK01JA0004	JDM	Weak FSNS	140 kDa band *
UK01ZE0005	JDM overlap	Homogeneous	Ku
UK01MB0006	JDM	FSNS	155/140 kDa bands *
UK01SP0007	JDM	Weak FSNS	155/140 kDa bands *
UK01ZB0008	JDM	Nucleolar	PM-Scl
UK01MV0010	JDM overlap	Weak Homogeneous with Nucleolar	Topoisomerase
UK01LP0016	JDM	Weak FSNS	Negative
UK01AP0017	JDM	Negative	140 kDa band *
UK01JC0018	JDM	Weak FSNS	155/140 kDa bands *
UK01SO0019	JDM	Negative	Negative
UK01SM0020	JDM overlap	Nucleolar	PM-Scl
UK01NT0021	JDM	Negative	Negative
UK01RK0022	JDM overlap	DCNS with Cytoplasmic Speckle	Jo-1
UK01RC0023	JDM	Negative	140 kDa band *
UK01KS0024	JDM	FSNS	Mi-2
UK01SP0025	JDM	Weak FSNS	155/140 kDa bands *
UK01CM0026	JDM	Negative	Unknown bands
UK01IP0027	JDM	DCNS	Unknown bands
UK01AP0028	JDM	Negative	Negative
UK01ES0029	JDM	Weak FSNS	140 kDa band *
UK01ZM0030	JDM overlap	Nucleolar	PM-Scl
UK01JB0031	JDM	Weak FSNS	Unknown bands
UK01JB0032	JDM	Negative	140 kDa band *
UK01FS0033	JDM	Weak FSNS	155/140 kDa bands *
UK01LT0037	JDM	Weak DCNS	Unknown bands
UK01RW0038	JDM	Negative	Unknown bands
UK01LS0039	JDM	DCNS	Unknown bands
UK01AL0040	JDM overlap	Nucleolar	PM-Scl
UK01EM0042	JDM	DCNS	Unknown bands

UK01KH0043	JDM	Homogeneous	Mi-2
UK01CB0047	JDM	Negative	Negative
UK01FU0048	JDM	Homogeneous	Mi-2
UK01GG0049	JDM	Negative	140 kDa band *
UK01JW0050	JDM overlap	Nucleolar with Coarse Speckle	PM-Scl
UK01LW0051	JDM overlap	Nucleolar with Cytoplasmic Speckle	PM-Scl
UK01LP0052	JDM	Negative	Negative
UK01EB0053	JDM	FSNS with Cytoplasmic Speckle	155/140 kDa bands *
UK01RS0054	JDM	FSNS	155/140 kDa bands *
UK01ST0056	JDM	Weak Fine Cytoplasmic Speckle	Negative
UK01EB0057	JDM	FSNS	155/140 kDa bands *
UK01VA0058	JDM	Homogeneous	155/140 kDa bands *
UK01JG0059	JDM	DCNS	Unknown bands
UK01JR0060	JDM	Weak FSNS	155/140 kDa bands *
UK01EK0063	JDM	Weak FSNS	Negative
UK01LC0065	JDM overlap	Nucleolar with Weak Homogeneous	U3-RNP
UK02NS0066	JDM	FSNS	140 kDa band *
UK01KS0068	JDM	Negative	140 kDa band *
UK01JJ0069	JDM	Weak FSNS	155/140 kDa bands *
UK01PB0070	JDM	Negative	140 kDa band *
UK01SB0071	JDM	Weak FSNS	140 kDa band *
UK01BH0073	JDM	Weak FSNS	Unknown bands
UK01CW0074	JDM	FSNS	Mi-2
UK01NW0075	JDM	FSNS	Unknown bands
UK01EB0076	JDM	FSNS	155/140 kDa bands *
UK01JT0077	JDM	Homogeneous	Mi-2
UK08KW0079	JDM	Weak FSNS	Unknown bands
UK04LM0082	JDM overlap	DCNS	U1-RNP
UK04JC0084	JDM	Weak DCNS	Negative
UK04BF0085	JDM overlap	DCNS	U1-RNP
UK04VL0086	JDM overlap	DCNS	U1-RNP
UK10CS0091	JDM	Negative	140 kDa band *
UK05AM0093	JDM	Negative	140 kDa band *
UK04NC0094	JDM	DCNS	Unknown bands



UK04GF0095	JDM overlap	Nucleolar with Discrete Speckle	PM-Scl
UK10KM0096	JDM	Homogeneous	Mi-2
UK09TH0098	JDM	Weak DCNS	140 kDa band *
UK09LW0099	JDM	FSNS	155/140 kDa bands *
UK09CD0101	JDM	DCNS	Negative
UK09SA0104	JDM	Weak Homogeneous	140 kDa band *
UK09HB0105	JDM	Negative	140 kDa band *
UK09LM0110	JDM	FSNS	155/140 kDa bands *
UK06RW0115	JDM	Weak Homogeneous	155/140 kDa bands *
UK01LT0118	JDM	$\gamma$	Unknown bands
UK02HC0119	JDM	FSNS	155/140 kDa bands *
UK01EB0120	JDM	Weak Homogeneous	140 kDa band *
UK10AW0122	JDM	DCNS	U1-RNP
UK01KS0125	JDM overlap	Nucleolar with Cytoplasmic Speckle	Jo-1
UK07AO0126	JDM	Weak DCNS	155/140 kDa bands *
UK07LF0128	JDM	$\gamma$	Negative
UK07ZM0129	JDM	Weak FSNS	140 kDa band *
UK07MG0130	JDM	FSNS	155/140 kDa bands *
UK07SL0131	JDM	DCNS	140 kDa band *
UK07MP0132	JDM	Weak Cytoskeletal	Unknown bands
UK08IC0135	JDM	Negative	140 kDa band *
UK08CT0136	JDM	FSNS	Unknown bands
UK08KP0137	JDM	Weak Cytoskeletal	Negative
UK08AD0138	JDM	Negative	Negative
UK10SW0139	JDM	DCNS	140 kDa band *
UK01LT0140	JDM	Homogeneous	Mi-2
UK01SD0141	JDM	FSNS	Unknown bands
UK01ET0142	JDM	DCNS	155/140 kDa bands *
UK01NZ0144	JDM	DCNS	Mi-2
UK07LH0148	JDM	Weak FSNS	140 kDa band *
UK08PS0149	JDM	Homogeneous	Mi-2
UK10NE0152	JDM	Negative	140 kDa band *
UK07AH0154	JDM	Weak Cytoplasmic Speckle	Negative
UK10LS0156	JDM	Weak FSNS	Negative
UK01BT0157	JDM	FSNS	155/140 kDa bands *

UK01KN0158	JDM	Weak FSNS	155/140 kDa bands *
UK01LG0161	JDM	FSNS	155/140 kDa bands *
UK01AP0162	JDM	Negative	Negative
UK01CH0164	JDM	Weak FSNS	155/140 kDa bands *
UK01AW0166	JDM	Weak FSNS	140 kDa band *
UK01AW0167	JDM	Negative	Negative
UK01LR0169	JDM	Negative	140 kDa band *
UK01AV0171	JDM overlap	DCNS	U1-RNP
UK01SB0172	JDM	Negative	140 kDa band *
UK09HB0174	JDM	DCNS	U1-RNP
UK01EC0175	JDM	$\gamma$	140 kDa band *
UK01IA0176	JDM	Weak Cytoplasmic Speckle	140 kDa band *
UK01JA0177	JDM overlap	DCNS	U1-RNP
UK01JS0178	JDM	Negative	140 kDa band *
UK04CJ0179	JDM	$\gamma$	140 kDa band *
UK01TB0180	JDM	$\gamma$	Negative
UK01PO182	JDM	DCNS	U1-RNP
UK01LF0183	JDM	Weak Homogeneous with	Negative
UK01SA0185	JDM	Nucleolar	PM-Scl
UK01NN0186	JDM	DCNS	Unknown bands
UK01CH0188	JDM	Weak FSNS	Unknown bands
UK01LF0189	JDM	Homogeneous	Mi-2
UK01GW0190	JDM	Negative	Unknown bands
UK01CL0192	JDM	Negative	Negative
UK07LG0194	JDM	$\gamma$	155/140 kDa bands *
UK01EW0196	JDM	Weak Cytoskeletal	Negative
UK01ST0197	JDM	$\gamma$	Negative
UK04JN0198	JDM	$\gamma$	155/140 kDa bands *
UK07RG0199	JDM	$\gamma$	140 kDa band *
UK10CK0200	JDM	$\gamma$	Negative
UK01AA0201	JDM	$\gamma$	140 kDa band *
UK04HS0203	JDM overlap	Nucleolar	PM-Scl
UK01AD0204	JDM	$\gamma$	Negative
UK01CA0206	JDM	$\gamma$	Negative
UK10CK0208	JDM	$\gamma$	Unknown bands
UK08LG0209	JDM	$\gamma$	140 kDa band *

UK01JH0212	JDM	γ	Negative
UK01TM0213	JDM	γ	Negative
UK01RH0217	JDM overlap	DCNS	U1-RNP
UK01AL0220	JDM	γ	155/140 kDa bands *
UK01BP0221	JDM	γ	Unknown bands
UK01TA0222	JDM	γ	140 kDa band *
UK09NR0223	JDM	γ	155/140 kDa bands *
UK01AS0224	JDM	γ	140 kDa band *
UK01RW0225	JDM	γ	140 kDa band *
UK01LB0226	JDM	γ	Negative
UK09AM0227	JDM overlap	γ	Unknown bands
UK01KS0228	JDM	γ	140 kDa band *
UK01GM0229	JDM overlap	Nucleolar	PM-Scl
UK01EF0231	JDM	γ	Negative
UK01AJ0233	JDM	γ	Negative
UK01OW0234	JDM	γ	Mi-2
UK01CW0236	JDM overlap	γ	Unknown bands
UK01SF0237	JDM	γ	Negative
UK01PB0239	Indeterminate	γ	Negative
UK10YK0244	JDM	γ	Negative
UK01RM0246	JDM	γ	Negative
UK01RG0247	JDM	γ	Negative
UK10KS0248	JDM	γ	PM-Scl
UK04CS0250	JDM	γ	Unknown bands
UK01KQ0251	Indeterminate	γ	Unknown bands

FSNS – fine speckle nucleolar sparing

DCNS – diffuse coarse nucleolar sparing

γ - ANA IIF not performed (insufficient serum available)

## **5.5 Appendix III**

### **Acknowledgements**

I would like to thank Mrs J Dunphy and Mrs P Owen for their assistance and guidance in performing ANA IIF on the juvenile myositis samples. Thank you to Dr Zoe Betteridge for her guidance with my immunoprecipitation studies. I would also like to thank Prof Lucy Wedderburn, Scientific Director for the Juvenile Dermatomyositis UK Cohort Study for her collaboration, guidance and support. On behalf of the Juvenile Dermatomyositis Research Group I would like to thank all the patients and their families who contributed to the Juvenile Dermatomyositis Cohort Study. I thank all local research coordinators and principal investigators who have made this research would not have been possible. The members who contributed were: Mr Ian Roberts, The Royal Liverpool Children's Hospital, Alder Hey, Liverpool, and Booth Hall Children's Hospital, Manchester; Dr Eileen Baildam, Booth Hall Children's Hospital, Manchester and now Alder Hey, Liverpool,; Dr Phil Riley, Booth Hall Children's Hospital, Manchester; Mrs Janis Scott and Dr Clive Ryder, Birmingham Children's Hospital, Birmingham; Mrs Gillian Jackson and Dr Sue Wyatt, Leeds General Infirmary, Leeds; Ms Elizabeth Camp and Dr Janet Gardner-Medwin, The Royal Hospital for Sick Children, Yorkhill, Glasgow; Mrs Alison Swift, Dr Helen Foster and Dr Mark Friswell, The Royal Victoria Infirmary, Newcastle; Mrs Elizabeth Hutchinson and Dr Helen Venning, Queens Medical Centre, Nottingham and Dr Clarissa Pilkington, Dr N Hasson and Ms Sue Maillard, Great Ormond Street Hospital, London. I thank H Varsani for technical assistance, in particular preparation of the serum samples, which were sent to the RNHRD for this study. The Arthritis Research Campaign awarded me the Barbara Ansell fellowship in Paediatric Rheumatology (grant 18136) to perform this work. I am grateful to the Arthritis Research Campaign who funded the autoantibody testing for this study as part of my fellowship.

## CHAPTER SIX

### RESULTS

#### Identification and characterisation of novel autoantigen systems in juvenile dermatomyositis

##### 6.1 Introduction

Classifying patients using a clinico-serological approach may lead to the identification of more homogeneous subsets within the JDM spectrum and therefore have prognostic implications.

The work described in Chapter 5 demonstrates that around 20% of children with JDM or JDM-overlap have well-defined autoantibody specificities (anti-ARS, anti-Mi-2 and MAAs). Therefore, approximately 80% of children would normally be serologically classified as either being non-specifically ANA positive or in some cases seronegative, when tested by ANA IIF or standard ENA ELISA. Over the last few years, a number of novel autoantibodies have been described in adult IIM (as outlined in Chapters 3 and 4). Similarly, there has been a preliminary report from Oddis *et al* of a new autoantibody termed anti-MJ, which targets a ~140-142 kDa protein in JDM (222). A further autoantigen target with a molecular weight of 155 kDa (reported in most cases as a doublet protein with a second weaker band at 140 kDa) has been described in JDM in a study by Targoff *et al* (143). The same novel specificity anti-p155/140 is detected in adult DM, which was described in the same study (143) and was also seen in the RNHRD adult IIM study (see Chapter 4). Two further studies have described anti-p155/140 in separate adult DM cohorts (157, 158). As previously discussed in Chapter 4, anti-p155/140 in adult DM is associated with more severe skin disease but more significantly has a strong association with malignancy. In the Targoff *et al* study, anti-p155/140 was seen in approximately 30% of their JDM population, however the clinical specificity of this cohort was not described (143).

The purpose of the work described in this chapter was to establish the frequency and to define the clinical significance of anti-p155/140 and anti-p140 autoantibodies in children recruited to the UK JDM Cohort Study. The secondary aims were to confirm whether the same p155/140 autoantigen is targeted in adult DM and to demonstrate p155/140 and p140 are different targets. As presented in Chapter 5, 40% of patients tested by IPP were found to be positive for these two different

autoantibodies. Both autoantibodies appear to define distinct clinical phenotypes within the JDM spectrum. This work has been published in the following journals:

- Gunawardena H, Wedderburn LR, North J, Betteridge Z, Dunphy J, Chinoy H, Davidson JE, Cooper RG, McHugh NJ; Juvenile Dermatomyositis Research Group UK. Clinical associations of autoantibodies to a p155/140 kDa doublet protein in juvenile dermatomyositis. *Rheumatology*. 2007;47:324-8
- Gunawardena H, Wedderburn LR, Chinoy H, Betteridge ZE, North J, Ollier WE, Cooper RG, Oddis CV, Ramanan AV, Davidson JE, McHugh NJ; Juvenile Dermatomyositis Research Group, UK and Ireland. Autoantibodies to a 140-kd protein in juvenile dermatomyositis are associated with calcinosis. *Arthritis Rheum*. 2009;60:1807-14.

## **6.2 Patients and Methods**

### **6.2.1 Patients and sera**

Subjects for the study described in this chapter were recruited from:

- The Juvenile Dermatomyositis Registry and Repository, UK and Ireland (24).

Full details are given in Chapter 5 (see Section 5.2.1). In brief, the JDRR has recruited patients from 10 centres around the UK (for details see appendix (24)) with juvenile-onset myositis, below the age of 16 years at disease onset and diagnosis, to form the JDM UK Cohort Study. All JDM cases had probable or definite disease according to *Bohan and Peter* criteria (1, 2). Demographic and clinical data using a standardised proforma were recorded at diagnosis and prospectively at subsequent visits, on average every 6 months (see Chapter 2, Section 2.1.2 and 2.1.3, and Appendix). Data were stored using anonymous codes onto a central database. Serum samples were taken at the time of diagnosis and stored at -20°C until required.

#### **6.2.1.1 Study 1 (Anti-p155/140 study)**

The first 116 patients recruited to the JDM National Registry and Repository (UK and Ireland) (now termed the JDM UK Cohort Study) were initially studied. The median age at symptom onset was 6 years (IQR 3, 9) and at diagnosis was 7 years (IQR 4, 10). The median follow-up from disease onset to time of this study was 69.5 months (IQR 47.5, 105.2) and 44.2 months (IQR 35.7, 65.8) from date of entry into

the registry to time of this study. Clinical information included the degree of skin involvement; muscle strength tested by the Childhood Myositis Assessment Scale (CMAS) (see Chapter 5, Appendix) (188), Physician's 10-point global assessment (PGA) (visual analogue scale) and serum muscle enzymes. When evaluating muscle involvement, only those JDM patients with documented muscle enzymes at disease onset / diagnosis (with subsequent maximum CK or LDH during follow up) were included for analysis. Similarly, those with a baseline CMAS or PGA at the time of entry to the registry (plus subsequent serial measures) were included. The type of skin lesion, in particular skin ulceration, oedema and Gottron's lesions combined with the distribution (extent) of skin involvement plus CMAS/ PGA score was defined as a marker of disease severity for the purpose of this study.

#### **6.2.1.2 Study 2 (Anti-p140 study)**

##### ***Patients***

Serum samples for serological typing were available from the overall cohort of 162 children recruited to the registry (including the first 116 children initially recruited / analysed in the anti-p155/140 study). Clinical data were available on 160 children and 74% were female. The median age of disease onset was 6 years, inter-quartile range (IQR) 3-9 and median age at diagnosis was 7 years (IQR 4, 10). The median follow-up from disease onset to the time of data analysis for this study was 48 months (IQR 33, 72) for the overall cohort. One hundred and thirty seven children had JDM. JDM-scleroderma overlap (JDM-SSc) is well recognised in JDM children with a history of Raynaud's phenomenon, sclerodacty and other sclerodermatous skin changes. In this study, 21 children were defined as JDM-SSc with two or more of the above features. Two children were defined as having other forms of juvenile myositis, not specifically JDM or JDM-SSc.

##### ***Controls***

Sera from 124 juvenile disease controls; twenty scleroderma / linear scleroderma, 8 systemic lupus erythematosus and 96 juvenile idiopathic arthritis cases were also analysed. No sera from healthy children were available, reflecting ethical difficulties in studies of this nature. Therefore, sera from 50 healthy adult controls were also serotyped.

## **6.2.2 Serological methods**

### ***Study 1 and 2***

#### ***Indirect immunofluorescence (IIF)***

IIF was performed by standard methods using HEp-2 cells and fluorescein-labelled anti-human IgG immunoglobulin (Sigma, UK).

#### ***Protein Radio-Immunoprecipitation (IPP)***

IPP from K562 cell extracts was performed as previously described in detail in Chapter 2. Briefly, 10 µl sera was mixed with 2 mg protein-A-Sepharose beads (Sigma, UK) in IPP buffer (10 mM Tris-Cl pH 8.0, 500 mM NaCl, 0.1% v/v Igepal) at room temperature for 30 min. Beads were washed in IPP buffer prior to the addition of 120 µl [<sup>35</sup>S] methionine labelled K562 cell extract. Samples were mixed at 4°C for 2 hr. Beads were washed in IPP buffer followed by TBS buffer (10 mM Tris-Cl pH 7.4, 150 mM NaCl) before being resuspended in 50 µl SDS sample buffer (Sigma, UK). After heating, proteins were fractionated by 10% SDS PAGE, enhanced, fixed and dried. Labelled proteins were analysed by autoradiography.

#### ***Immunodepletion experiments***

##### ***Study 1***

Immunodepletion experiments were performed using reference anti-p155/140-JDM-positive sera and reference anti-p155/140-adult-positive sera (adult sera from the RHHRD serology study – see Chapter 4, Section 4.3.4), to ascertain if the IPP pattern seen was due to precipitation of the same p155/140 autoantigen. Cell extracts were depleted of autoantibody targets using anti-p155/140-JDM-positive serum or anti-p155/140-adult DM-positive serum and normal serum as a negative control. These extracts were then used in further immunoprecipitations using both juvenile and adult anti-p155/140 positive serum.

As described in Chapter 2, Section 2.3.4: duplicate samples each containing 10 mg protein A Sepharose beads in 1 ml IPP buffer and 50 µl reference anti-p155/140 serum were mixed with end-over-end rotation at room temperature for 30 min. Beads were washed four times in 1 ml IPP buffer and 1 tube (A) was placed on ice whilst 150 µl [<sup>35</sup>S] methionine-labelled K562 cell extract and 350 µl IPP buffer was added to the remaining tube (B). Tube B was mixed at 4°C for 2 hr after which the supernatant was transferred to tube A, this was mixed at 4°C for a further 2 hr. The supernatant from tube A was then transferred to a fresh tube (C) and stored at -80°C. IPP using JDM or adult DM serum and either 150µl control [<sup>35</sup>S]-methionine-



labelled cell extract or the immunodepleted supernatants (C) were completed as described for IPP using [<sup>35</sup>S] methionine.

## **Study 2**

### ***Immunodepletion experiments with anti-p140-JDM and anti-p155/140-JDM sera***

Immunodepletion experiments were performed using reference anti-140-JDM-positive sera and reference anti-p155/140-JDM-positive sera to confirm that these autoantibodies target different autoantigens. Cell extracts were depleted of autoantibody targets using anti-140-JDM positive serum or anti-p155/140-JDM positive serum and normal serum as a negative control. These extracts were then used in further immunoprecipitations using different JDM anti-p140 and anti-p155/140 positive serum. As described in Chapter 2, Section 2.3.4: duplicate samples each containing 10 mg protein A Sepharose beads in 1 ml IPP buffer and 50 µl reference anti-p140 or reference anti-p155/140 serum were mixed with end-over-end rotation at room temperature for 30 min. Beads were washed four times in 1 ml IPP buffer and 1 tube (A) was placed on ice whilst 150 µl [<sup>35</sup>S] methionine-labelled K562 cell extract and 350 µl IPP buffer was added to the remaining tube (B). Tube B was mixed at 4°C for 2 hr after which the supernatant was transferred to tube A, this was mixed at 4°C for a further 2 hr. The supernatant from tube A was then transferred to a fresh tube (C) and stored at -80°C. IPP using JDM serum and either 150µl control [<sup>35</sup>S]-methionine-labelled cell extract or the immunodepleted supernatants (C) were completed as described for IPP using [<sup>35</sup>S] methionine.

### ***IPP with mouse monoclonal anti-NXP-2 and immunodepletion experiments***

Preliminary data from Oddis *et al* and Targoff *et al* have described autoantibodies to the MJ antigen, which is a 140-142 kDa protein in JDM (222, 228). The MJ target has recently been identified as nuclear matrix protein NXP-2, also termed MORC-3 (228). Immunodepletion experiments were performed to investigate the identity of the p140 target in this UK JDM cohort, which is likely to be the same as the previously identified MJ autoantigen. Therefore, immunodepletion experiments were performed using different reference anti-p140-JDM-positive sera and a commercial mouse monoclonal anti-NXP-2 antibody. Ten µl of anti-p140 positive sera or 50 µl of commercial mouse antibody to NXP-2 (MORC3) (Medical & Biological Laboratories, Nagoya, Japan) were mixed with 100 µl of prewashed protein G Dynabeads (Dyna, Liverpool, UK) in sodium phosphate (ph 8.1, 0.1 M) at room temperature for 30 min. The antigens were immunoprecipitated as described for IPP, using [<sup>35</sup>S] methionine labelled K562 cell extract. Immunodepletion was performed to ascertain whether the

IPP pattern observed with anti-p140 positive JDM sera and the commercial antibody to NXP-2 were due to precipitation of the same antigen. Cell extracts were depleted of autoantibody targets using a reference anti-p140 positive JDM sera and normal serum (NS) as a negative control. In brief, duplicate samples each containing 10 mg protein A sepharose beads (when preparing pre-depleted p140 cell extract for IPP with commercial anti-NXP-2, 150 µl prewashed protein G Dynabeads were used) in 1 ml IPP buffer and 50 µl anti-p140 serum (or 50 µl anti-p155/140 serum) were mixed at room temperature for 30 min. The beads were washed in IPP buffer and 1 tube (A) was placed on ice whilst 120 µl [<sup>35</sup>S] methionine labelled K562 cell extract and 380 µl IPP buffer was added to tube (B). Tube B was mixed at 4°C for 2 h, the supernatant was then transferred to the corresponding tube A, which was then mixed at 4°C for a further 2 h. The supernatant from the corresponding tubes (A) i.e. p140 antigen depleted cell extract were stored at -80°C. IPP with depleted cell extracts were completed using 50 µl commercial anti-NXP-2, and different anti-p140 sera (10 µl).

#### ***IPP Western blotting with mouse monoclonal anti-NXP-2***

As well as immunodepletion experiments, immunoprecipitation western blotting studies using mouse monoclonal anti-NXP-2 antibodies were also performed. Forty µl of reference sera (anti-p140-JDM sera, anti-p155/140-JDM sera, anti-Mi-2-JDM sera) and 40 µl of normal sera were incubated with washed Sepharose protein A beads. Samples were centrifuged and the supernatant removed. Samples were washed twice with 1 ml triethanolamine and the supernatant removed. Samples were cross-linked and incubated with 1 ml 5mM BS<sup>3</sup> in triethanolamine (2.86 mg / ml – 25 mg into 8.74 ml), centrifuged and the supernatant removed. Samples were further incubated with 1 ml 50mM Tris-Cl and then washed three times with PBS and twice with IPP buffer prior to the addition of 1 ml unlabelled K562 cell extract. Samples were mixed at 4°C for 2 hr. Beads were washed in IPP buffer followed by TBS buffer (10 mM Tris-Cl pH 7.4, 150 mM NaCl) before being resuspended in 80 µl SDS sample buffer (Sigma, UK). After heating, proteins were fractionated by 10% SDS PAGE.

Nitrocellulose membrane and blotting paper was soaked in Transfer Buffer (Trizma Base, sodium acetate, EDTA, 0.1% SDS and 20% methanol). A sandwich of blotting paper, gel, nitrocellulose and blotting paper was placed in a transfer cage. Proteins were transferred to nitrocellulose membrane following electrophoresis. The nitrocellulose membrane was washed with Ponceau stain, rinsed with water and air-

dried overnight. The membrane was washed with 0.05% PBS-Tween followed by blocking solution (10% milk powder in PBS-Tween – 10 grm in 100 ml, pH 7.2). The membrane was then incubated with primary antibody (commercial mouse monoclonal anti-NXP-2 antibody – 1:200 dilution – 50 µl in 10 ml block solution) for 1 hr. The membrane was then washed in PBS-Tween, and then incubated with secondary anti-mouse IgG antibody (AbCam, UK), at 1:10,000 dilution in block solution – 50 µl in 50 ml, pH 7.2, for 1 hr. The membrane was washed again with PBS-Tween and water, and then air-dried.

### **6.2.3 Statistical analysis**

The frequencies of clinical features were compared using the Chi-squared test with Yates' continuity correction or the Fisher's exact test for groups with small numbers. Where data was not normally distributed the Mann-Whitney-U test was used to compare continuous data. Median values (inter-quartile ranges) were expressed where appropriate and  $P$  values  $<0.05$  were considered significant.  $P$  values ( $P_{\text{corr}}$ ) were adjusted using the Bonferroni correction. SPSS for Windows (version 14) (for clinical data) was used to perform statistical analysis.

## **6.3 Results**

### **6.3.1 Anti-p155/140: autoantibodies targeting a 155/140 kDa doublet protein in juvenile dermatomyositis**

#### ***Serological results***

Following IPP, sera from a number of JDM patients recognised two distinct proteins forming a doublet with molecular weights of 155 kDa and 140 kDa (see Figure 16). The same pattern was observed in a subset of adult DM patients (see Chapter 4, Figure 13 and this chapter, Figure 16). Non-specific weak nuclear patterns were observed on IIF between anti-p155/140 patients (data not shown).

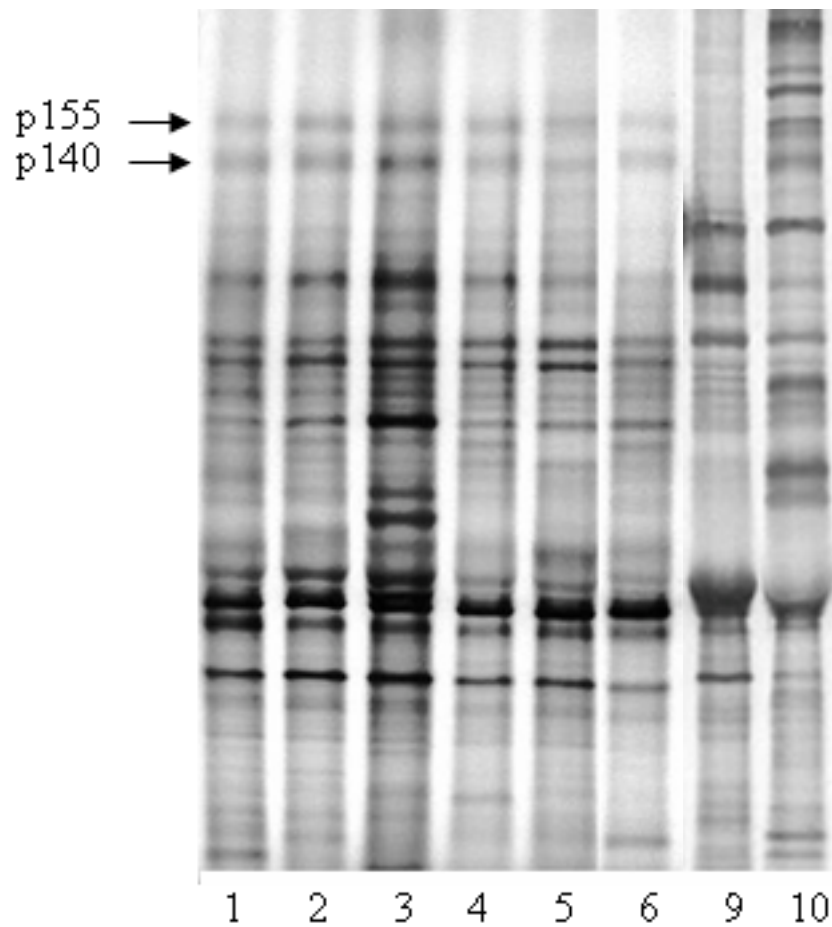
The immunodepletion results support the co-identity of the p155/140 kDa doublet precipitated by sera from both JDM and adult DM groups (see Figure 17). When the cell extracts were pre-depleted with normal serum, no targets were removed from the extract and the 155 kDa and 140 kDa autoantigens were still precipitated by juvenile and adult sera. However, when the cell extract was pre-depleted with either juvenile or adult anti-p155/140 positive sera, the autoantigens were no longer detectable in juvenile or adult anti-p155/140 positive sera respectively. This provided

good evidence that the sera from JDM and adult DM contained the same autoantibody specificity.

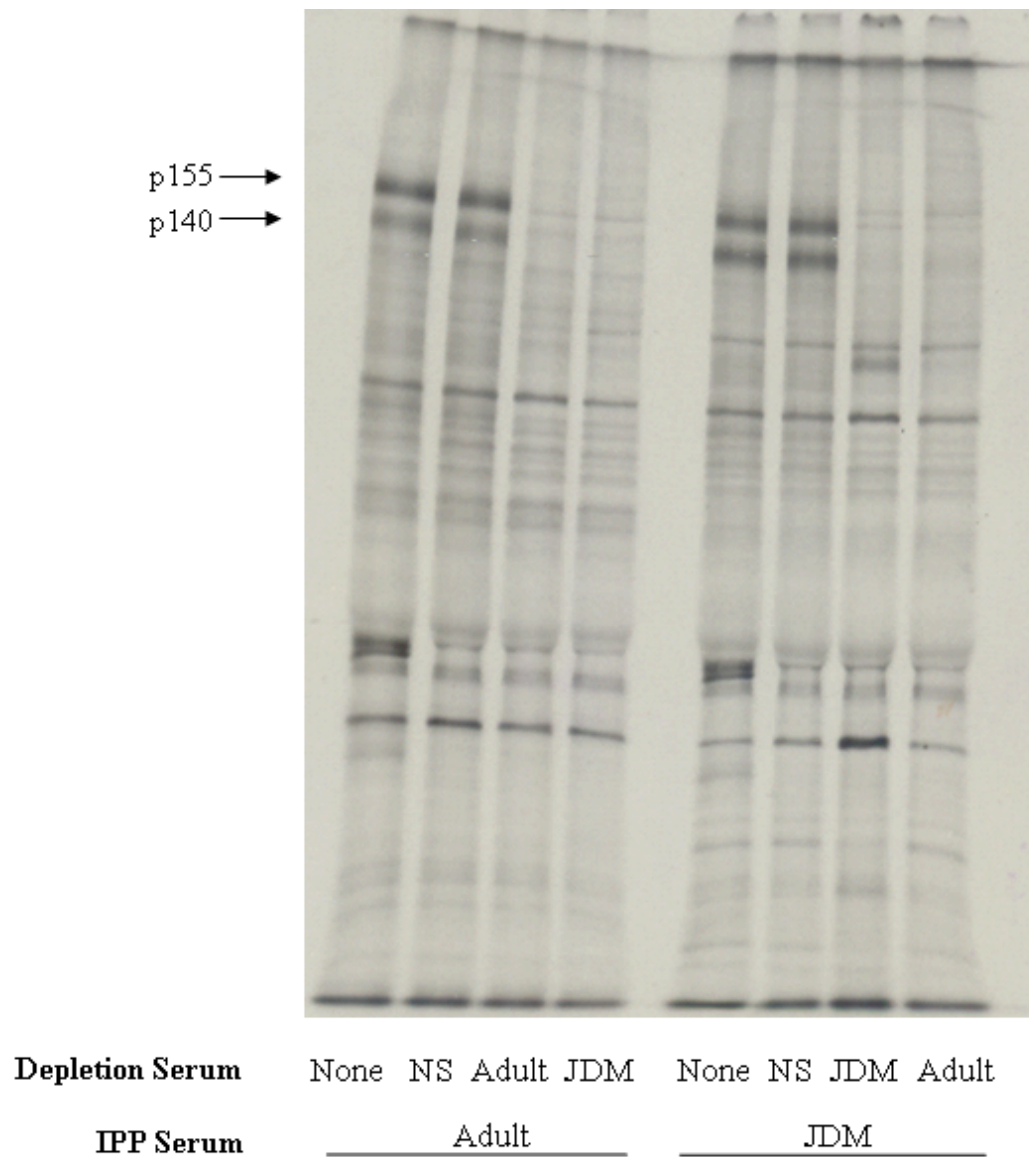
***Frequency and clinical features of anti-p155/140 autoantibodies***

From 116 juvenile myositis sera, 27 (23%) had anti-p155/140 autoantibodies. Information on the degree of skin involvement and other selected clinical features are outlined in Table 18. There was a higher frequency of males in the anti-p155/140 positive children compared to anti-p155/140 negative children ( $P_{\text{corr}}=0.04$ ). Anti-p155/140 positive JDM patients had an increased frequency of skin lesions (Gottron's papules  $P_{\text{corr}}=0.027$ , ulceration  $P_{\text{corr}}=0.045$ ) with a wider distribution of cutaneous involvement, particularly over the small joints ( $P_{\text{corr}}<0.001$ ) and large joints ( $P_{\text{corr}}=0.027$ ). Cutaneous oedema was also more frequent in anti-p155/140-positive children ( $P=0.013$ ) but this was not significant following multiple corrections. Overall, there was no significant difference in those with elevated muscle enzymes at diagnosis or during disease course or those with an abnormal MRI or muscle biopsy between anti-p155/140 positive and negative groups (data not shown). However not all children had data on this, in particular some did not have a MRI or biopsy performed. There was a trend towards lower CMAS (lower values indicate more severe weakness) and higher PGA in anti-p155/140 positive children at baseline and during follow up, although this did not reach statistical significance. The frequency of other clinical signs including lipoatrophy, arthritis, Raynaud's, sclerodermatous skin changes, dysphagia, mouth ulcers or alopecia was not significantly different between children with or without anti-p155/140. There was no history of malignancy in the entire JDM cohort during the follow-up period.

**Figure 16:** Immunoprecipitation of p155/140 kDa autoantigens. 10% SDS PAGE of immunoprecipitates of [<sup>35</sup>S] labelled K562 cell extract. Sera used for immunoprecipitation include Lanes 1-3; adult anti-p155/140 positive serum, Lanes 4-6; juvenile anti-p155/140 positive serum, Lane 9; juvenile anti-Jo-1 (histidyl tRNA synthetase) positive serum, Lane 10; juvenile anti-Mi-2 positive serum. Positions of the p155 and p140 antigens are indicated on the left.



**Figure 17:** Immunodepletion experiments - Autoradiogram of 10% SDS-PAGE of immunoprecipitates using either anti-p155/140-adult DM sera or anti-p155/140-JDM positive sera. Immunoprecipitation was performed with control [<sup>35</sup>S]-methionine-labelled cell extract or [<sup>35</sup>S]-methionine-labelled cell extract depleted with either normal serum (NS), adult positive anti-p155/140 serum or JDM positive anti-p155/140 serum. The bands corresponding to the p155 and p140 autoantigens are indicated.



**Table 18: Selected clinical associations of anti-p155/140 autoantibodies in JDM patients**

	Anti-p155/140		<i>P</i> <sub>corr</sub>
	Positive (n=27)	Negative (n=89)	
Age at diagnosis, median (IQR)	6 years (4, 10)	7 years (5, 10)	
Male:Female	44.4	20.2	0.04
<b>Type of skin lesion</b>			
Gotttron's papules	100	76.1 <sup>#</sup>	0.027
Ulceration	51.9	21.3	0.045
Oedema	63	33.7	<i>NS</i> *
Calcinosis	11.1	23.6	<i>NS</i>
Lipoatrophy	17.9	14.3	<i>NS</i>
<b>Distribution of skin lesion</b>			
Periorbital	92.6	65.9 <sup>#</sup>	<i>NS</i>
Periungal	81.5	59.1 <sup>#</sup>	<i>NS</i>
Trunk	6 (22.2)	10 <sup>#</sup> (11.4)	<i>NS</i>
Small joints	100	62.5 <sup>#</sup>	<0.001
Large joints	77.8	43.2 <sup>#</sup>	0.027
<b>Muscle disease<sup>‡</sup></b>			
Baseline CMAS, median (IQR)	36 (13.8, 48)	44 (35, 50.5)	<i>NS</i>
Lowest CMAS, median (IQR)	36 (13.8, 46.8)	43 (29, 49.0)	<i>NS</i>
Baseline PGA, median (IQR)	4.7 (2.0, 7.1)	2.8 (1.1, 5.0)	<i>NS</i>
Highest PGA, median (IQR)	5.5 (2.5, 7.1)	3 (1.2, 5.1)	<i>NS</i>

Values in percentages unless otherwise indicated. <sup>#</sup>Data on 88 children. <sup>‡</sup>Not all patients had data available for each clinical feature. CMAS (childhood myositis assessment scale: 0-53) at baseline and lowest (worst) score during follow-up. PGA (physician's global assessment: 0-10) at baseline and highest (worst) score during follow-up.

\**P*<sub>(uncorr)</sub>=0.013

### **6.3.2 Anti-p140: autoantibodies targeting a nuclear matrix protein NXP-2 in juvenile dermatomyositis**

#### ***Serological results***

##### ***Identification of anti-p140 autoantibodies in JDM sera***

Following IPP, sera from a number of JDM patients recognized a distinct protein band with a molecular weight of ~140 kDa (see Figure 18). No anti-p140 positive sera were found to immunoprecipitate any other known MSAs or MAAs. A weak non-specific nuclear pattern or in some cases a negative ANA was observed on IIF in all anti-p140 sera. Anti-p140 was not detected in any of the juvenile disease control sera or healthy adult control sera.

##### ***Immunodepletion experiments using anti-140 and anti-p155/140 JDM sera***

Immunodepletion confirmed anti-p140 sera target a different protein to JDM sera that recognises the p155/140 doublet protein (see Figure 19, lanes 1-4) showing respective bands are still present following IPP).

The IPP Western blotting with mouse monoclonal anti-NXP-2 studies were negative (did not yield any results). No reference p140 bands were recognised following incubation with primary mouse monoclonal antibody anti-NXP-2 and secondary anti-mouse IgG antibody. This may be because of loss of immunoreactivity of the p140 autoantigen following transfer from SDS-PAGE to nitrocellulose membrane.

##### ***Confirmation of the p140 autoantigen***

The results suggest that the p140 protein targeted by JDM sera in this study has the same identity as the MJ antigen, NXP-2 (228). Using a commercial antibody raised against NXP-2, IPP resulted in the precipitation of a band with the same molecular weight and IPP pattern as that observed in anti-p140 positive JDM sera (see Figure 20, lane 3). When cell extract was pre-depleted with JDM anti-p140 positive sera, the IPP band present with commercial anti-NXP-2 was no longer detectable (Figure 20, lane 4). The immunodepletion results supported the co-identity of the p140 protein precipitated by different anti-p140 positive juvenile sera. Figure 19 (lanes 5-6) shows an example where the 140 kDa band is no longer detectable following IPP with anti-p140 sera (1) and (2) using pre-depleted reference p140 cell extract.

##### ***Clinical associations of anti-p140 autoantibodies***

Of 162 juvenile myositis sera serotyped by IPP, 37 (23%) had anti-p140 autoantibodies. This autoantibody specificity was found exclusively in JDM patients,



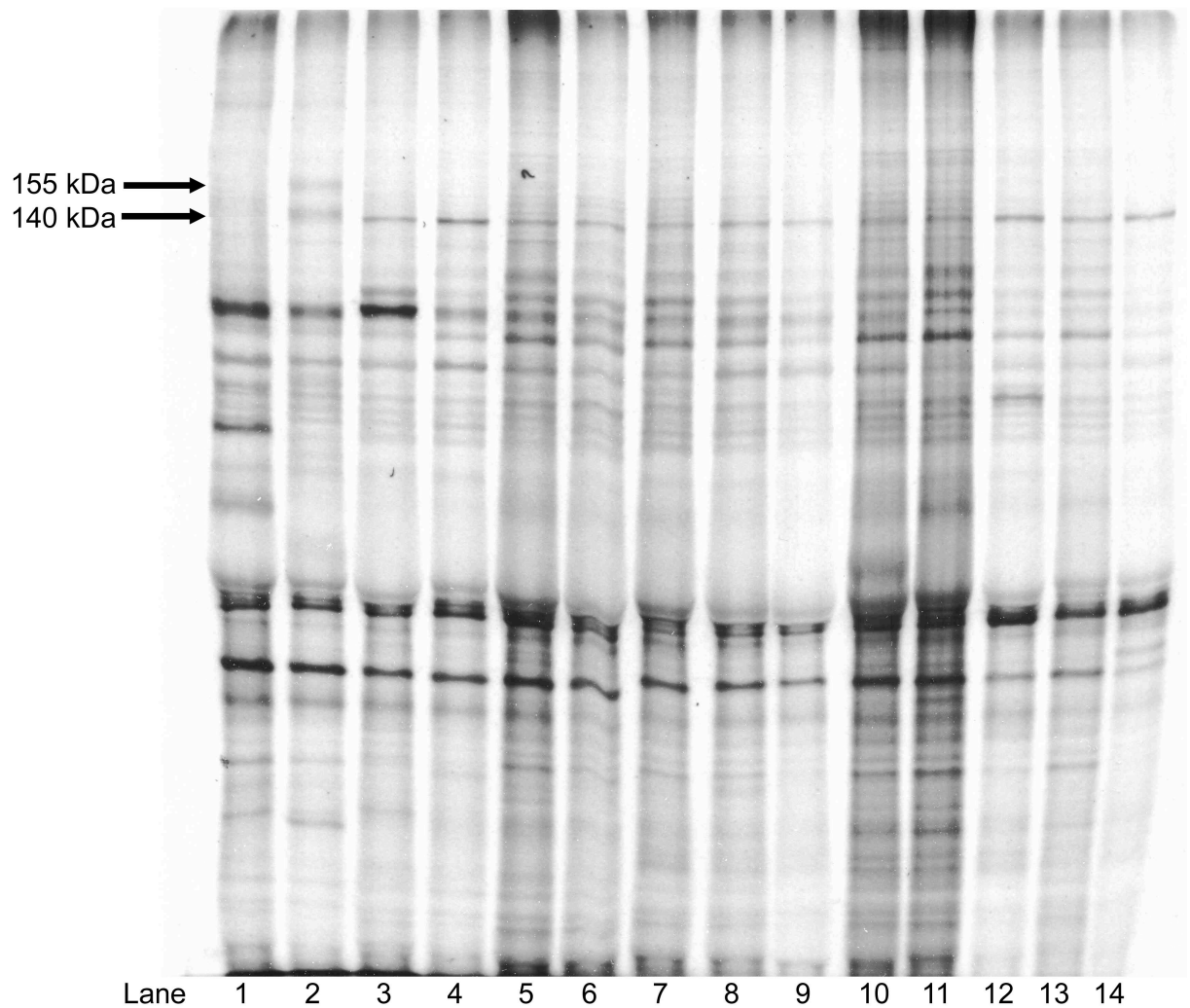
with a frequency of 27%. Anti-p140 was not detected in any overlap JDM-SSc sera. Selected clinical features of anti-p140 positive patients compared to the overall anti-p140 negative JDM cohort are shown in Table 19. Table 20 shows a comparison between anti-p140, anti-p155/140 and MAA positive cases. There was no significant difference in female:male ratio, age at onset or diagnosis and disease duration (from disease onset to time of this study) between anti-p140 positive and negative cases. Overall there was no significant difference in the presence of Gottron's lesions, skin ulceration and oedema or the distribution of rash except anti-p140 positive children had no rashes on the trunk compared to anti-p140 negatives ( $P_{\text{corr}}=0.02$ ). Anti-p140 positive cases had a significant association with the presence of subcutaneous calcinosis compared to anti-p140 negative patients ( $P_{\text{corr}}<0.005$ , odds ratio 7.0, 95% CI 3.0-16.1). There was no significant difference between other clinical features when comparing anti-p140 positives *versus* negatives including baseline CK, CMAS, childhood health assessment questionnaire (CHAQ), physician's global assessment scale (PGA), and the presence of arthritis, Raynaud's, dysphagia, mouth ulcers and alopecia (data not shown).

Possible clinical differences were noted when anti-p140 positive patients were compared to anti-p155/140 positive patients (Table 18). Overall, age of onset or diagnosis and disease duration (anti-p140 positives, median 48 months IQR 34, 72 *versus* anti-p155/140 positives, median 52 months, IQR 36, 84) was similar between the autoantibody groups. Compared with anti-p155/140 positive children, anti-p140 positives again had an association with calcinosis, 54% versus 14%,  $P_{\text{corr}}=0.015$  OR 7.1 (2-25). In contrast, anti-p155/140 positives compared with the anti-p140 positive group had a higher frequency of ulceration and cutaneous oedema however this result was not significant after correcting for multiple comparisons. The distribution of rash was wider on the trunk  $P_{\text{corr}}<0.005$  and over the small joints ( $P_{\text{corr}}=0.05$  OR 13.5 95% CI 2-113) in anti-p155/140 compared with anti-p140 positive cases. At the time of diagnosis, anti-p140 compared to anti-p155/140 positive children had a non-significantly lower CK (median 202, IQR 76, 2142 versus median 571 IQR 234, 2495) and LDH (median 845 IQR 710, 1620 versus median 1171 IQR 736, 1647). In addition, anti-p140 positives had a higher CMAS at diagnosis compared to anti-p155/140 positives (median 42, IQR 23, 49 versus median 16, IQR 7, 38) (not significant after adjustment for multiple comparisons). There was a non-significant trend towards a lower baseline CHAQ and PGA in the anti-p140 group compared to the anti-p155/140 group; median CHAQ 1.31 (0.75, 1.63) versus 1.63 (0.78, 2.34), median PGA 5 (3, 7.4) versus 7.3 (5, 7.8) respectively. However, this data needs to

be interpreted in the context that a small number of patients in either group had this data recorded at diagnosis.

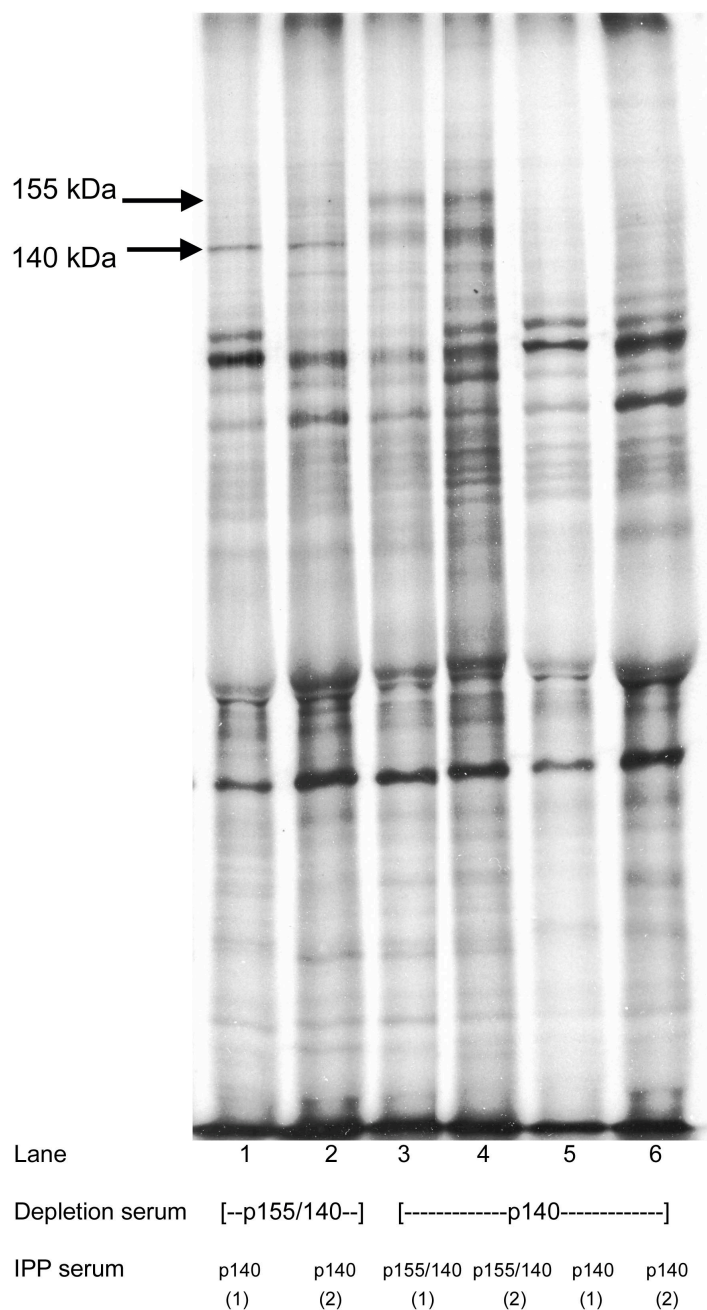
**Figure 18: Immunoprecipitation of p140 autoantigens**

Autoradiogram of 10% SDS PAGE of immunoprecipitates of [<sup>35</sup>S] labelled K562 cell extract. Sera used for immunoprecipitation include Lane 1; normal serum, Lane 2; anti-p155/140 positive JDM serum, Lanes 3-14; different anti-p140 positive JDM sera.



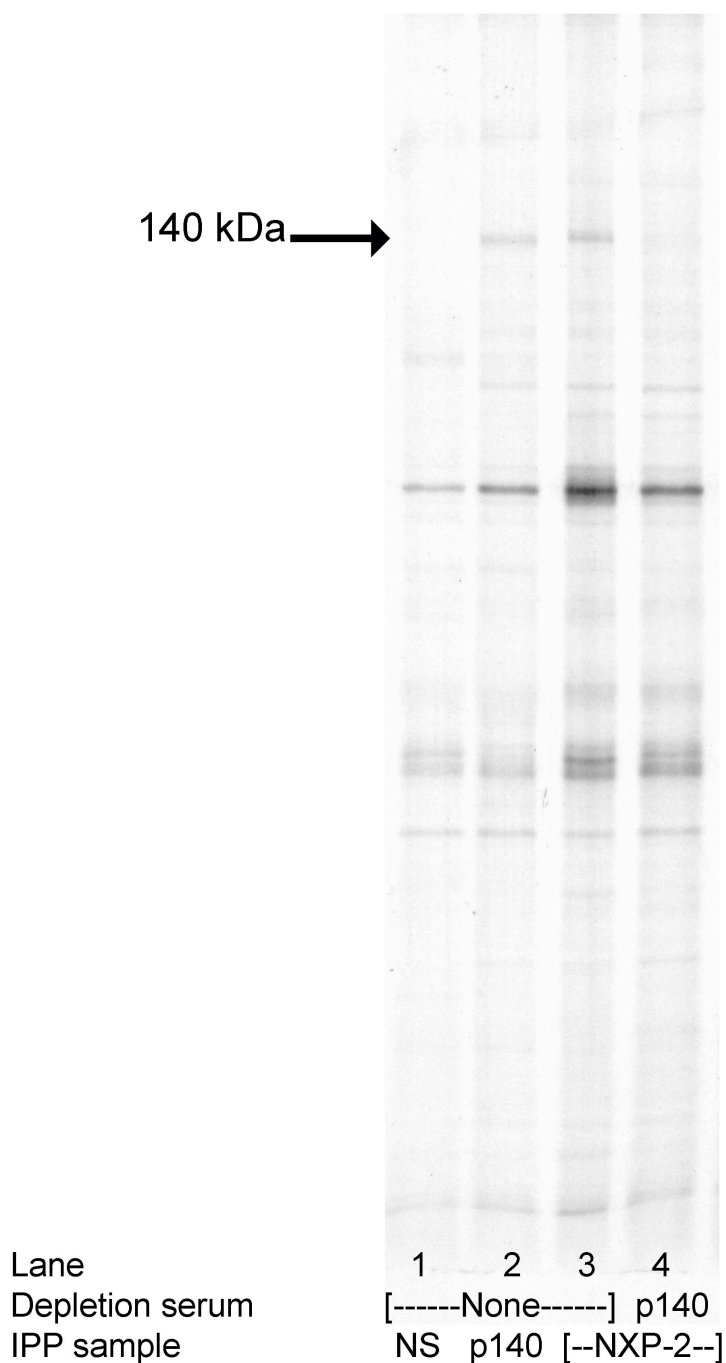
### Figure 19: Immunodepletion experiments with anti-p140 and anti-p155/140

Autoradiogram of 10% SDS-PAGE of immunoprecipitates using different anti-p140 positive JDM sera (lanes 1-2 and 5-6) and anti-p155/140 positive JDM sera (lanes 3-4). Immunoprecipitation was performed with [<sup>35</sup>S] labelled K562 cell extract pre-depleted with either reference anti-p140 positive JDM serum or reference anti-p155/140 positive JDM serum.



**Figure 20: Immunoprecipitation and immunodepletion of mouse monoclonal nuclear matrix protein NXP-2**

Autoradiogram of 10% SDS-PAGE of immunoprecipitates of [<sup>35</sup>S] labelled K562 cell extract using normal sera (NS) (lane 1), reference anti-p140 positive JDM sera (lane 2) and commercial anti-NXP-2 (lane 3). Lane 4, immunoprecipitation using commercial anti-NXP-2 with [<sup>35</sup>S] labelled K562 cell extract pre-depleted with reference anti-p140 positive JDM serum.



**Table 19: Selected clinical associations of anti-p140 positive JDM patients compared with JDM patients overall (anti-p140 negative) and with anti-p155/140 positive JDM patients \***

	Anti-p140 positive (n=37) **	All JDM anti- p140 negative (n=125)**	Anti-p155/140 positive (n=28)
Age at onset (yrs)	6	6	6
Median (IQR)	(2, 10)	(4, 9)	(4, 10)
Age at diagnosis (yrs)	7	7	7
Median (IQR)	(4.5, 10.3)	(4, 10)	(4, 10)
Female	72	74	57
<b>Type of skin lesion</b>			
Gottron's papules	85	82	100
Ulceration	34	23	57
Oedema	35	35	64
Calcinosis	54 ¶	15	14
Lipoatrophy	18	13	18
<b>Distribution of skin rash</b>			
Periorbital	79	69	96
Periungal	64	67	86
Small joints	67 #	72	96
Large joints	52	51	75
Trunk	0 ##	18	32

\* Values are the percentage of patients, unless otherwise stated.

\*\* Not all patients had the clinical data available for each feature.

$P_{\text{corr}}$ : corrected  $P$  values, OR: odds ratio (with 95% confidence intervals).

¶  $P_{\text{corr}} < 0.005$ , OR 7.0 (3.0 - 16.1) versus all JDM anti-p140 negative patients and  $P_{\text{corr}} = 0.015$ , OR 7.1 (2-25) versus anti-p155/140 positive patients.

#  $P_{\text{corr}} = 0.05$  OR 13.5 (2-113) anti-p155/140 positive patients versus anti-p140 positive patients.

##  $P_{\text{corr}} = 0.02$  versus all JDM anti-p140 negative patients and  $P_{\text{corr}} < 0.005$  versus anti-p155/140 positive patients.

**Table 20: Clinical associations of anti-p140-JDM patients compared with anti-p155/140 positive-JDM, and MAA-positive- JDM patients \***

	Anti-p140 positive (n=37) **	Anti-p155/140 positive (n=28) **	MAA- positive (n=23) **
Age at onset (yrs)	6	6	6
Median (IQR)	(2, 10)	(4, 10)	(4, 10)
Age at diagnosis (yrs)	7	7	9
Median (IQR)	(4.5, 10.3)	(4, 10)	(5, 11)
Female	72	57	78
<b>Type of skin lesion</b>			
Gottron's papules	85	100 ¶	65
Ulceration	34	57 ¶¶	13
Oedema	35	64 α	17
Calcinosis	54 αα	14	22
Lipoatrophy	18	18	35
<b>Distribution of skin rash</b>			
Periorbital	79	96	35
Periungal	64	86 λ λ	57
Small joints	67	96 #	65
Large joints	52	75 ##	30
Trunk	0	32 ###	17
<b>Other features</b>			
Arthritis	49	39	65
Sclerodermatous features	3	4	61 Ɔ
Raynaud's phenomenon	11	11	48 Ɔ Ɔ
Dysphagia	31	43	35
Mouth ulcers	29	32	4 γ
Alopecia	37	39	26

## Key for Table 20

\* Values are the percentage of patients, unless otherwise stated.

\*\* Not all patients had the clinical data available for each feature.

$P_{\text{corr}}$ : corrected  $P$  values

¶  $P_{\text{corr}}=0.01$ , anti-p155/140-positives versus MAA-positives

¶¶  $P_{\text{corr}}=0.03$ , anti-p155/140-positives versus MAA-positives

α  $P_{\text{corr}}=0.02$ , anti-p155/140-positives versus MAA-positives

αα  $P_{\text{corr}}=0.01$ , anti-p140-positives versus anti-p155/140-positives / MAA-positives

λ  $P_{\text{corr}} < 0.001$ , anti-p155/140-positives versus MAA-positives

#  $P_{\text{corr}}=0.045$ , anti-p155/140-positives versus MAA-positives / anti-p140-positives

##  $P_{\text{corr}}=0.03$ , anti-p155/140-positives versus MAA-positives

###  $P_{\text{corr}}=0.01$ , anti-p155/140-positives versus MAA-positives and anti-p140-positives

⊃  $P_{\text{corr}} < 0.001$ , MAA-positives versus anti-p140 / anti-p155/140-positives

⊃⊃  $P_{\text{corr}} < 0.001$ , MAA-positives versus anti-p140 / anti-p155/140-positives

λ λ  $P_{\text{corr}}=\text{NS}$  ( $P=0.045$ , anti-p155/140-positives versus MAA-positives / anti-p140-positives)

γ  $P_{\text{corr}}=\text{NS}$  ( $P=0.04$ , anti-p155/140 and anti-p140-positives versus MAA-positives)



## 6.4 Discussion

As described in this thesis and in previous studies there is now increasing evidence that MSAs are associated with clinical subsets in the adult IIM spectrum. The work described in Chapter 5 and 6 suggests serological profiles may also be associated with clinical phenotypes in JDM. Previously, myositis autoantibodies were detected infrequently in JDM with anti-Mi-2 having the strongest association (see Chapter 5) (125, 127). Therefore, detection and characterisation of novel MSAs in JDM may identify distinct clinical subsets within this disease group. Identification of new autoimmune markers may help clinicians predict clinical outcomes and lead to further insights in disease pathogenesis.

There have been reports of novel protein targets including p140 kDa and p155 kDa polypeptides in both adult and juvenile DM. Two reports have described autoantibodies targeting a 140 kDa protein (anti-MJ) and a 155/(140) kDa protein in US JDM patients (143, 222). In adult DM, anti-p155/140 autoantibodies are associated with a history of malignancy (see Chapter 4) (143, 157, 158). In the study described in Chapter 4 (see Section 4.3.4, Table 12), anti-p155/140 autoantibodies were also detected in adults with CADM with no history of cancer and although the numbers are small, this group of adults tended to be younger. In this cohort of children with JDM, no malignancy has been reported to date in those with anti-p155/140 autoantibodies; long term follow up studies will be required to ascertain whether there is any association with malignancy later in life in this group. In addition, validation of anti-p155/140 in larger groups of both adult and juvenile DM is required. An important finding of this study is the demonstration by immunodepletion that anti-p155/140 autoantibodies appear to target the same autoantigen in both adult and juvenile DM. Targoff *et al* described anti-p155 (in most cases with a second weaker 140 kDa band) in approximately 30% of their JDM population (143). The clinical specificity of this cohort was not described. This study describes detailed clinical features of JDM patients with autoantibodies targeting p155/140 and confirms they have similar cutaneous clinical associations to adults with the same autoantibody specificity (as described in Chapter 4, Section 4.3.4, Table 11). Like adult DM cases with anti-p155/140 these children have more extensive skin involvement including Gottron's papules over a wider distribution. Anti-p155/140 autoantibodies also appear to define a subset of JDM with more peripheral oedema and skin ulceration. There was a trend towards lower baseline and worst ever CMAS, and higher PGA in anti-p155/140 positive JDM compared to negative patients, although this was not statistically significant. In summary, anti-p155/140

autoantibodies occur frequently in JDM and the preliminary findings suggest that they may identify patients with more severe skin and possibly muscle disease. Identification of this autoantibody at diagnosis may help predict the clinical course.

In this chapter, a further novel autoantibody specificity in the JDM UK cohort is described. Anti-p140 autoantibodies form a major serological subset in juvenile myositis found exclusively in JDM and not detected in any JDM-SSc overlap cases. Anti-p140 positive sera recognised the same polypeptide and did not immunoprecipitate any other known myositis autoantigens, including the anti-p155/140 autoantibody. Combining the data from Study 1 and 2 in this chapter, anti-p140 and anti-p155/140 autoantibodies are detected in approximately 40% of JDM cases, in contrast to a much lower frequency of anti-Mi-2 and other myositis-specific or associated autoantibodies as published previously (10, 125, 127, 219).

The confirmation of a further serological subset in JDM appears to have important clinical implications. Anti-p140 has a significant association with the presence of calcinosis when compared to the overall juvenile myositis cohort. In addition, anti-p140 and anti-p155/140 appear to define JDM into two serological subsets with more homogeneous clinical features. Combined with the data from Study 1 in this chapter, anti-p155/140 positive children appear to have a wider distribution of skin disease, more cutaneous complications including oedema or ulceration and possibly overall higher disease activity but a significantly lower frequency of calcinosis compared to anti-p140 positive children. The clinical differences observed between anti-p140, anti-p155/140 positives and patients without either of these specificities are not explained by differences in time between disease onset or disease duration. This is an interesting observation because factors suggested to influence the development of calcinosis are persistent active disease including chronic cutaneous inflammation (185, 201, 229). The association between serotype and clinical phenotype suggests that the targeted autoantigens p140 and p155/140 may play role in the pathogenesis of skin and soft tissue complications in JDM. Based on previous work that showed an association with the TNF $\alpha$ -308A allele, an increased production of tumour necrosis factor- $\alpha$  and calcinosis (201); future studies to investigate for other potential susceptibility genes including TNF polymorphisms in JDM patients with anti-p140 or anti p155/140 autoantibodies would be of major interest.

The p140 protein targeted by an autoimmune response in our JDM cohort study is consistent with nuclear matrix protein NXP-2, the MJ autoantigen described in preliminary reports in a US JDM cohort, and most recently in an Argentine JDM patient study by Espada *et al* (222, 225, 228). In the latter study, RNA and protein immunoprecipitation and immunoblotting tested 64 patients with juvenile IIM for autoantibodies. The anti-MJ autoantibody was detected exclusively in 25% of cases, a similar frequency to the anti-p140 autoantibody found in the UK JDM cohort study described here. Clinical features of anti-MJ autoantibodies suggest this specificity is a marker of disease severity, although the numbers were small and no direct comparisons with other serological groups was made. Muscle contractures (44%) and lipoatrophy (44%) were more frequent in anti-MJ-positive cases. This specificity was detected predominantly in pure JDM cases with proximal weakness and classic skin changes. The frequency of calcinosis was 31% and cutaneous vasculitis (with skin ulceration) 38% respectively. Clinical similarities with the anti-MJ and anti-p140 autoantibody are noted between this Argentinean cohort and the UK JDM cohort. The main differences, in particular the prevalence of lipoatrophy, may represent variations in clinical evaluation and length of follow-up for each study (225).

In the study by Espada *et al* 22% of juvenile patients (all JDM) were positive for the anti-p155/140 autoantibody, again a similar number to that detected in the work in this chapter. All children had proximal muscle weakness with a high prevalence of muscle contracture, and the majority has classic DM skin disease. There was also a relatively high frequency of cutaneous vasculitis with persistent ulceration (57%) (225), the same frequency noted in anti-p155/140-positive children from the UK cohort study (see Table 18 and 19).

A further clinically important complication of JDM is lipodystrophy or lipoatrophy (a loss of subcutaneous fat, which is either localised or generalised affecting the face, trunk and limbs). Previous studies have reported prevalence rates between 12-40% (24, 230, 231). In a recent study, Bingham *et al* reported a lower frequency of lipodystrophy of 8% in a tertiary referral centre juvenile IIM registry (232). In the Bingham *et al* study, generalised lipodystrophy was shown to be associated with the anti-p155 (anti-p155/140) autoantibody (143), as well a chronic disease course, muscle atrophy, joint contractures, and interestingly calcinosis (232). In this UK cohort study, the overall frequency of lipodystrophy / atrophy was 14% (see Chapter 5, Section 5.3.2, Table 16), although the prevalence of this complication in anti-p155/140 and anti-p140 positive children was not significantly different compared to

the rest of the cohort. Interestingly, lipoatrophy was seen in a relatively higher frequency of children who were positive for MAAs, however this did not reach statistical significance. A potential reason for the difference in lipoatrophy associations between this study and the work by Bingham *et al* (232) is the way respective clinicians define lipodystrophy or lipoatrophy. Furthermore, the relatively higher frequency of lipoatrophy in MAA-positive children may be because this overall autoantibody group was associated with JDM-scleroderma overlap.

Larger collaborative studies are required to evaluate further the clinical specificity of both anti-p155/140 and anti-p140 autoantibodies in juvenile myositis populations. Work to date and the work presented in this chapter suggest that these two specificities are very important autoantibody systems with pathogenic, clinical and prognostic implications. When grouped together these two serological subsets are positive in between 40-50% of JDM cases, and appear to define those children with a more severe disease course including an association with widespread skin lesions, skin ulceration, oedema, lipodystrophy, calcinosis and worse muscle disease with muscle contracture (222, 225, 232, 233) (and Chapter 6).

The identity of the p155 kDa and p140 kDa doublet protein requires investigation. It is likely that the p155/140 autoantigen identified in this work is the same as the p155/140 doublet proteins identified in recent studies (143, 157, 158), although immunodepletion or reverse IPP blotting studies with prototype sera are required for further confirmation. A preliminary report has identified the target of the anti-p155 autoantibodies as transcriptional intermediary factor 1-gamma (TIF1- $\gamma$ ) (159). This nuclear protein is a member of a novel family of transcriptional coregulator-encoding genes, which function in cell differentiation and development (181). As yet, it is unclear what relationship the p140 protein (as part of the p155/140 doublet) has to the identified p155 TIF1- $\gamma$  autoantigen i.e. is this a separate protein, which forms part of a multi-protein complex, or is it simply a cleaved or modified fragment of the same protein.

Augmented expression myositis-specific autoantigens in diseased muscle and cancer tissue suggests a new paradigm for pathogenesis of IIM (107, 137). It is not clear why a tumour-associated autoantigen in adult DM should be recognised in JDM, particularly because there appears to be no association malignancy. However, other clinical features including the degree of skin disease in anti-p155/140 positive children and adults are very similar. Longitudinal prospective studies are required to

investigate whether this autoantibody in juvenile disease persists into adulthood and if so are they clinically significant. Perhaps the observation that the same autoantigen system is expressed in younger DM adults without cancer as well JDM suggests that some perturbation of p155/140 expression in proliferating cells combined with a more efficient anti-cancer response by a younger immune system may be a unifying mechanism to explain why it is targeted by an autoimmune response in children and adults with DM.

The p140 autoantigen NXP-2 has nuclear matrix binding, RNA-binding, and coiled-coil domains that are structurally separated, which implicates a role in diverse nuclear functions including regulation of transcription and RNA metabolism (228, 234). The other dermatomyositis-specific autoantigen targets in children and adults; Mi-2 and p155/140 (p155 – transcription intermediary factor 1-gamma) (159, 181) are nuclear proteins that also mediate gene transcription. It is of further interest to note that autoantibodies to small ubiquitin-like modifier enzyme (SAE), which is involved in post-transcriptional modification termed sumoylation, have been described in adult dermatomyositis (see Chapter 4, Section 4.3.2) (160). It is interesting that anti-SAE autoantibodies were not detected in any sera from the UK JDM cohort (Chapters 5 and 6), particularly because NXP-2 has been shown to be a sumoylation target involved in transcriptional repression (235). However, this observation may suggest shared pathogenic mechanisms in both juvenile and adult dermatomyositis.

In conclusion, anti-p155/140 and anti-p140 are clinically important serological markers found in a high frequency of JDM patients that are associated several disease complications, which confers significant morbidity. In the future, routine testing of these novel autoantibodies at disease onset could have prognostic value and identify those children at risk of more severe disease, which may influence management. Furthermore, increasing our understanding of autoimmune targets and their relationship to clinical phenotype in juvenile myositis may provide further insight into pathogenic pathways, which in turn will stimulate new therapeutic approaches.

## **6.5 Appendix**

### **I. Acknowledgements**

I would like to thank Mrs J Dunphy and Mrs P Owen for their assistance and guidance in performing ANA IIF on the juvenile myositis samples. Thank you to Dr Zoe Betteridge for her guidance with my immunoprecipitation, immunodepletion and western blotting studies. I would also like to thank Prof Lucy Wedderburn, Scientific Director for the Juvenile Dermatomyositis UK Cohort Study for her collaboration, guidance and support.

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## **CHAPTER SEVEN**

### **DISCUSSION**

#### **Introduction**

This chapter will outline two preliminary studies conducted alongside the main body of this thesis. The chapter will close with a general discussion of the work conducted as part of this thesis, and its relevance to the current understanding of the relationship between pathogenesis, serological subsets and clinical phenotypes in adult and juvenile IIM. To conclude, hypothetical models of autoimmune dysregulation in distinct microenvironments and how this may lead to disease initiation and propagation in IIM, will be explored. Potential future work focusing on antigen expression at a cellular and tissue level, and the development of solid phase assays for commercial testing of novel myositis-specific autoantibodies (MSAs), will be discussed.

#### **7.1 Other preliminary studies**

##### **7.1.1 Autoantibodies in idiopathic interstitial pneumonia**

###### ***Introduction***

As described in previous chapters, interstitial pneumonia is a frequent and significant manifestation of adult IIM, particularly the anti-synthetase syndrome (ASS) subset. Moreover, interstitial lung disease may be the presenting or predominant manifestation of IIM, especially in ASS “sine myositis” and clinically-amyopathic DM (29, 31, 40-42).

NSIP has been reported to be the most frequent subtype on both HRCT and lung biopsy in IIM (38). AIP, OP and UIP have also been reported and identified as patterns responsible for acute or rapidly progressive interstitial pneumonia with AIP and UIP associated with the worst prognosis (28-31, 33, 39). It is well recognised that interstitial lung involvement may be the first or predominant organ manifestation preceding muscle or skin disease, particularly in the anti-synthetase syndrome (see Section 1.5.1) and in patients with clinically-amyopathic DM (CADM) (see Section 1.5.3). Similarly, it is now recognised that interstitial pneumonia can be the major complication of systemic sclerosis sine scleroderma, particularly in association with the specific scleroderma-associated autoantibody anti-Th/To (227, 236-238).

Diffuse parenchymal lung disease is labelled idiopathic interstitial pneumonia when there is no evidence of a causative agent or coexisting connective tissue disease. The standard autoimmune profile is usually negative, even in patients who later

manifest a connective tissue disease (CTD). Therefore some patients with interstitial pneumonia may have underlying CTD characterised by the presence of specific autoantibodies not detected on routine serological testing, particularly ANA ELISA (most commonly used in routine clinical laboratories). In this pilot study, a small cohort of patients previously diagnosed as having idiopathic interstitial pneumonia have been screened for CTD-associated autoantibodies using ANA IIF and protein immunoprecipitation (IPP).

### ***Patients and Methods***

Twenty patients with a previous clinico-radiological diagnosis of idiopathic interstitial pneumonia were studied (patients from the Royal United Hospital Interstitial Lung disease clinic were invited to participate and not pre-selected). On further detailed clinical review by a Rheumatologist (HG), some patients had clinical features suggestive of an underlying CTD including a history of Raynaud's phenomenon (see Table 22).

All patients had high-resolution computerised tomography (HRCT) scans, which were re-reviewed (see Section 3.2.2). The subtype of interstitial pneumonia was categorised based on the American Thoracic Society / European Respiratory Society classification (32).

Patients' sera were analysed using ANA indirect immunofluorescence (IIF) and further characterised using protein immunoprecipitation of <sup>35</sup>S-labelled K562 cells combined with SDS-PAGE to identify specific autoantigen targets, as described previously (see Chapter 2, Section 2.3.1 and 2.3.2).

### ***Results***

For detailed results see Table 21 and see Figure 21. Six patients had a positive ANA on IIF (>1/40 titre), all with a homogeneous pattern. Of those, two patients' sera were negative on IPP, three cases had strong unidentified bands, and one patient was anti-topoisomerase positive on IPP (characteristic 100 kDa band, and confirmed by ENA ELISA). Despite two patients testing negative on ANA IIF, both had unidentified bands on IPP.

Five patients' sera with negative anti-nuclear staining but a strong cytoplasmic speckle tested positive for recognised anti-aminoacyl tRNA synthetase (ARS)



autoantibodies (one anti-Jo-1, one anti-PL-7, two anti-PL-12, and one patient with possible anti-Ha autoantibodies). See Table 22.

### **Discussion**

In this preliminary study, 30% of patients with a previous diagnosis of idiopathic interstitial pneumonia actually tested positive for CTD-associated autoantibodies. One case tested positive for anti-topoisomerase autoantibodies, with NSIP lung disease, and clinical features of Raynaud's phenomenon. Therefore, the patient may be re-diagnosed as having systemic sclerosis sine scleroderma (major SSc organ involvement, in this case fibrotic lung disease, a positive SSc-autoantibody, without characteristic skin changes of scleroderma) (239). In two recent studies by Fischer *et al*, a subset of patients with idiopathic interstitial pneumonia had nucleolar staining patterns of ANA IIF, the majority of which were positive for anti-Th/To autoantibodies (236, 237). Anti-Th/To is associated with less sclerodermatous features and a higher risk of pulmonary hypertension and lung fibrosis (227, 239), and this autoantibody specificity should be considered in patients with these clinical manifestations.

Of those 30% with CTD-autoantibodies, the other cases were positive anti-ARS autoantibodies, in particular non-Jo-1-anti-ARS. All non-Jo-1-anti-ARS has a clinico-radiological pattern consistent with UIP, except the anti-PL-7 case, which had NSIP. As described in Chapter 3 (see Section 3.3.3), lung fibrosis is often the predominant clinical manifestation in non-Jo-1-ARS-positive patients. Furthermore, radiological patterns demonstrate more architectural distortion with pronounced traction bronchiectasis and honeycombing, and histologically fibroblastic foci. This subtype of interstitial pneumonia is synonymous with idiopathic pulmonary fibrosis (formerly termed cryptogenic fibrosing alveolitis). Therefore, it is possible a proportion of patients with idiopathic pulmonary fibrosis actually have non-Jo-1-ARS, especially anti-PL-12 autoantibodies (42, 150). Anti-PL-7 autoantibodies have also been shown to be significantly associated with interstitial pneumonia and much milder or subclinical muscle disease (145).

In summary, patients with a presumed idiopathic interstitial pneumonia may have subtle features of underlying CTD, and a negative routine autoimmune panel should not dissuade clinicians to consider more novel serological markers of autoimmunity in some. Moreover, even in patients with UIP lung disease, a negative ANA but a strong cytoplasmic pattern on IIF should prompt more detailed investigation. A

proportion of these patients may be a “*formes fruste*” of an underlying CTD, in particular anti-synthetase syndrome where lung disease is the only or predominant manifestation. Historically, the general consensus has been patients with idiopathic lung fibrosis have a limited response to immunomodulatory therapy, whereas patients with CTD-associated lung disease do appear to respond or stabilise with corticosteroid and immunosuppressive therapies. Thus, early identification of this group of patients will influence treatment strategies and long-term prognosis.

**Part of this preliminary study has been published in abstract form:**

Gunawardena H, Betteridge Z, Owen P, McHugh NJ, Foley NM. Novel markers of connective tissue disease in patients with idiopathic interstitial pneumonia. European Respiratory Journal 2008;32:320s.

**Table 21: Serological results in idiopathic interstitial pneumonia patients**

<b>Patient</b>	<b>Age</b>	<b>IIP subset</b>	<b>ANA on IIF</b>	<b>IPP</b>
R23840	61	NSIP	1/160 homogeneous	Negative
R23839	73	NSIP	Negative	Non-specific bands,
R23838	74	NSIP	Negative with strong cytoplasmic speckle	Anti-Jo-1
R23837	75	UIP	1/40 homogeneous	Negative
R23843	66	Indeterminate	Negative	Negative
R23842	46	NSIP	Negative	Negative
R24103	61	UIP	Negative	Unknown 112 kDa band
R24101	58	NSIP	Negative but strong cytoplasmic speckle	Anti-PL-7
R24102	53	RBILD	1/40 homogeneous and fine cytoplasmic speckle	Unknown 72 kDa band
R24100	58	NSIP	Negative	Negative
R25095	68	UIP	1/2560 homogeneous	Unknown 25 kDa band
R23038	62	UIP	Negative with strong cytoplasmic speckle	Anti-PL-12
R24526	59	Indeterminate	Negative with cytoplasmic speckle	Negative
R19338	65	UIP	1/640 homogeneous and fine cytoplasmic speckle	Unknown 72 kDa band
R25133	60	NSIP	Negative	Unknown 30 kDa band
R26806	36	UIP	Negative with strong cytoplasmic speckle	Anti-PL-12
R26812	51	UIP	Negative with strong cytoplasmic speckle	Anti-PL-12
R26518	64	NSIP	Homogeneous 1/640	Anti-Topoisomerase
R26895	58	UIP	Negative with weak fine cytoplasmic speckle	Unknown 31, 32 and 145 kDa bands
R15264	75	UIP	Negative with strong cytoplasmic speckle	Unknown 62-63 kDa band (Ha antigen?) Possible synthetase (not confirmed)

**Table 22: Clinical features of patients with presumed idiopathic interstitial pneumonia with recognised CTD-autoantibodies tested by IPP**

Patient	Clinical Feature					
	RP	MH	Fever	Arthralgia	Subtype of IP	Autoantibody
R26806	Yes	Yes		Yes	UIP	Anti-PL-12
R26518	Yes				NSIP	Anti-Topo
R15264	Yes	Yes	Yes		UIP	Anti-p62 (Anti-Ha?)
R24101		Yes			NSIP	Anti-PL-7
R23038			Yes	Yes	NSIP	Anti-Jo-1
R26812				Yes	UIP	Anti-PL-12

RP: Raynaud's phenomenon

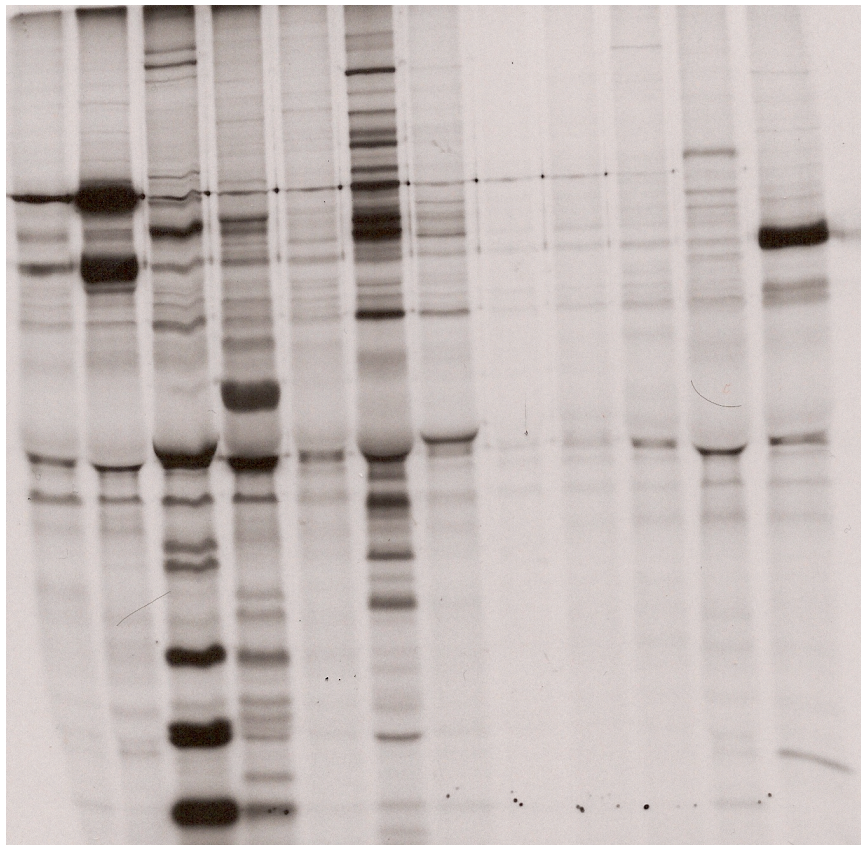
MH: Mechanic's hands

Topo: Topoisomerase

## Figure 21: Immunoprecipitation study in idiopathic lung disease patients

### 21A

Autoradiogram of 10% SDS-PAGE of immunoprecipitates of [<sup>35</sup>S] labelled K562 cell extract using normal serum (lane 1), reference anti-PL-12 and reference anti-PL-7 sera (lane 2), reference anti-Jo-1, reference anti-U1-RNP and reference anti-RNAP-II sera (lane 3), reference anti-PM-Scl, reference anti-Ro and reference anti-La sera (lane 4), R23840 serum (lane 5), R23839 serum (lane 6), R23838 (anti-Jo-1) serum (lane 7), R23837 serum (lane 8), R23843 serum (lane 9), R23842 serum (lane 10), R24103 serum (lane 11), and R24101 (anti-PL-7) serum (lane 12 – arrow).

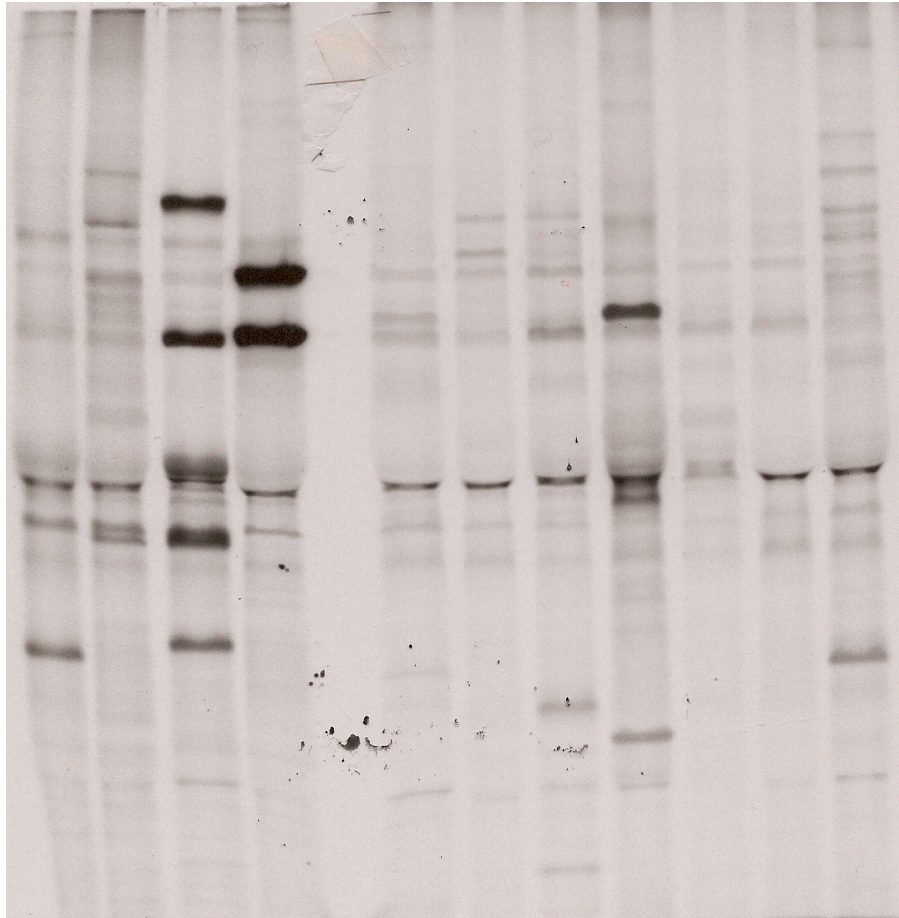


1 2 3 4 5 6 7 8 9 10 11 12

**Figure 21: Immunoprecipitation study in idiopathic lung disease patients**

**21B**

Autoradiogram of 10% SDS-PAGE of immunoprecipitates of [ $^{35}$ S] labelled K562 cell extract using reference anti-U3-RNP serum (lane 1), R2518 serum (anti-Topoisomerase) (lane 2), reference anti-mitochondrial serum (lane 3), reference anti-Ku serum (lane 4), R24102 serum (lane 5), R24100 serum (lane 6), R19803 serum (lane 7), R19338 serum (lane 8), control normal serum (lane 9), R24526 serum (lane 10), and R25133 serum (lane 11). Arrow – 100 kDa.

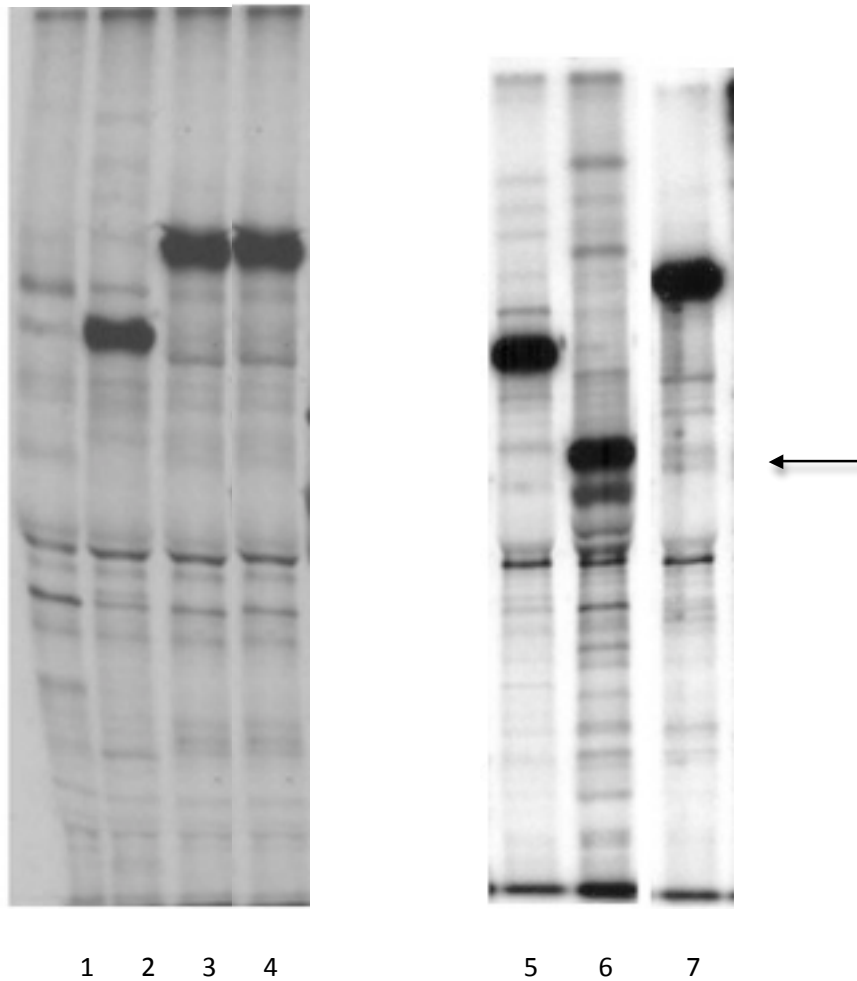


1 2 3 4 5 6 7 8 9 10 11

**Figure 21: Immunoprecipitation study in idiopathic lung disease patients**

**21C**

Autoradiogram of 10% SDS-PAGE of immunoprecipitates of [ $^{35}$ S] labelled K562 cell extract using control normal serum (lane 1), reference anti-PL-7 serum (lane 2), R26806 serum (anti-PL-12) (lane 3), R26812 serum (anti-PL-12) (lane 4), reference anti-PL-7 serum (lane 5), R15264 (anti-p62 / Ha) serum (lane 6) (arrow), and reference anti-PL-12 serum (lane 7).



### **7.1.2 Anti-p140 autoantibodies in adult IIM**

#### ***Introduction***

As described in Chapter 6, a major autoimmune target in JDM is the p140 autoantigen, likely to be the nuclear matrix protein NXP-2. Following detailed serological analysis of adult IIM patients recruited to the RNHRD cohort study and AOMIC UK Registry (work described in Chapters 3 and 4), a subset were noted to also immunoprecipitate a p140 kDa band. Therefore, preliminary work was undertaken to investigate if autoantibodies to the same p140 autoantigen, are present in adult myositis sera.

#### ***Patients and Methods***

Clinical data and serum samples were available from the UK Adult Onset Myositis Immunogenetic Collaboration (AOMIC) and the RNHRD IIM cohort study (as described in Chapters 2-4). As previously described, serum samples were autoantibody typed by immunoprecipitation using [<sup>35</sup>S] labelled K562 cells. Immunofluorescence using HEp-2 cells and fluorescein-labelled anti-human immunoglobulin was completed on adult anti-p140 positive samples.

#### ***Results***

Sera from 266 myositis patients were screened for anti-p140 autoantibodies. Overall, seven cases (3%) were positive for anti-p140 autoantibodies, which were detected exclusively in 5% of DM patients. From the RNHRD cohort study, two reference sera (R18883 and R23389) were positive for autoantibodies to a p140 kDa band. A further five patients' sera were found to recognise a 140 kDa band on SDS-PAGE. No anti-p140 antibody positive patients were positive for other recognized autoantibodies (see Figure 22). IPP experiments using cell extract pre-depleted with reference adult-p140-positive sera (R18883) suggest that the identity of the p140 target in adult DM may be the same as the p140 protein recognised in JDM, and different to the p155/140 autoantigen (see Figure 23). ANA IIF results are shown in Table 23. The major clinical features of anti-p140-positive patients were heliotrope rash (73%), Gottron's lesions (82%), periungal erythema (91%) and systemic involvement including weight loss or fever (78%). In particular, five out of seven (71%) anti-p140-positive adult patients had interstitial pneumonia. There was no cancer-associated myositis in the adult anti-p140 positive subset. In contrast to anti-p140-positive JDM patients where calcinosis is a significant feature, this was only present in one patient.



**Dr Zoe Betteridge (and HG) have since conducted further work. This work has been published in abstract form:**

Betteridge ZE, Gunawardena H, Chinoy H, Vencovsky J, Allard S, Gordon PA, Cooper RG, McHugh NJ. Autoantibodies to the p140 autoantigen NXP-2 in adult dermatomyositis. *Arthritis Rheum* 2009;60:S815.

In total, serum and clinical data has now been analysed in 443 adults with myositis (PM and DM), based on the Bohan and Peter diagnostic criteria, recruited to the Adult Onset Myositis Immunogenetic Collaboration, UK, RNHRD Cohort and the Institute of Rheumatology, Prague, CZ. All sera were autoantibody typed by immunoprecipitation using <sup>35</sup>S-labelled K562 cells. Immunofluorescence using Hep-2 cells and fluorescein-labelled anti-human immunoglobulin was completed on p140 positive samples.

Thirteen (2.93%) adult patients with IIM were positive for anti-p140 autoantibodies. All anti-p140 positive patients had DM. The prevalence of anti-p140 autoantibodies specifically in the DM cohort was 5.86%. Anti-p140 autoantibodies were not detected in sera from normal healthy controls or PM, myositis-overlap, SSc or SLE patients.

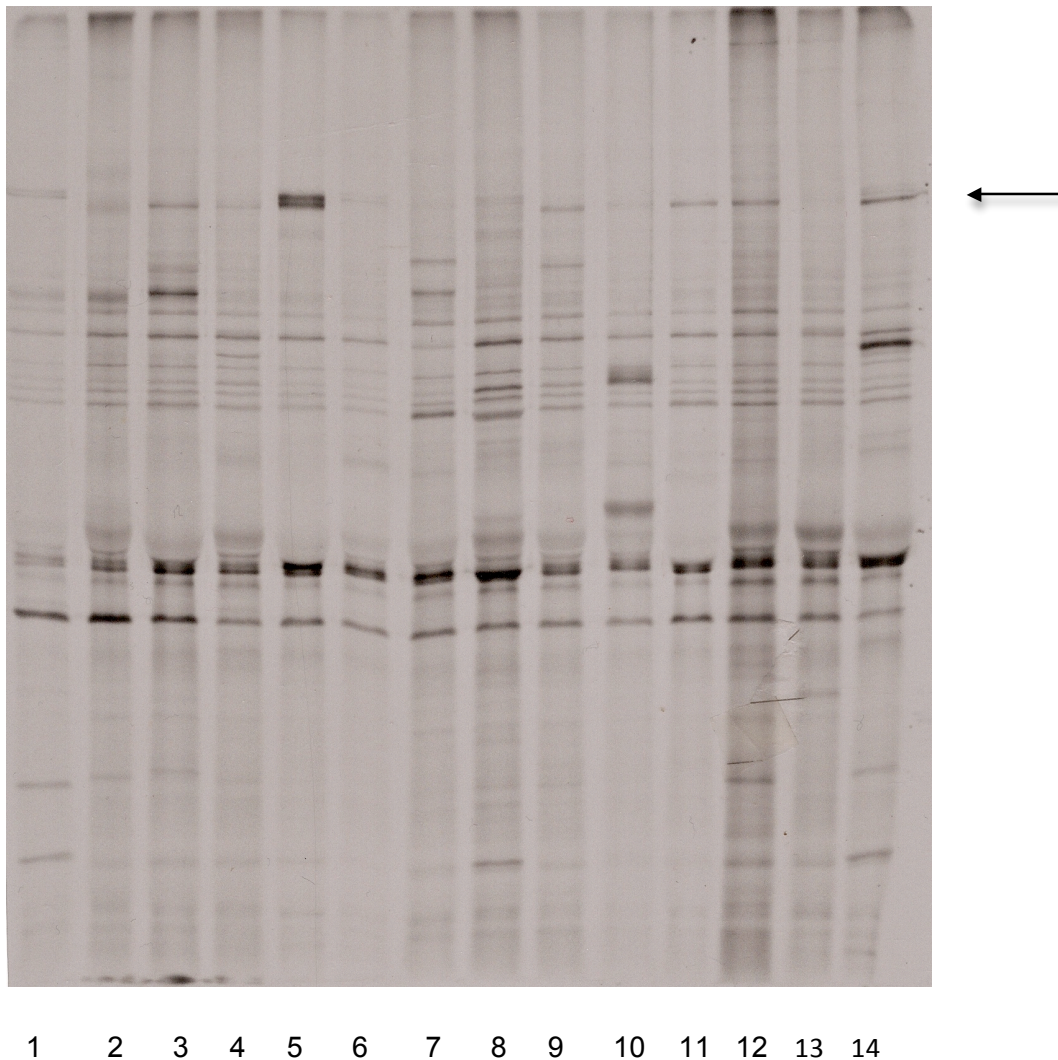
The major clinical associations of anti-p140 autoantibodies in adults are shown in Table 24. In particular, the frequency of interstitial pneumonia in anti-p140 positive patients was 61.5% in comparison to 26.2% in the DM anti-p140 negative group ( $p=0.016$ ). In addition, no p140 positive patients had cancer-associated myositis and in comparison with the JDM cohort, where calcinosis was a significant association, calcinosis was only present in two adult p140-positive patients (15.4%).

### ***Discussion***

Anti-p140 autoantibodies form a further serological subset in adult DM. In addition, preliminary data suggests that the clinical associations of anti-p140 autoantibodies in adults differ from JDM. In particular, lung disease appears to be a major feature of anti-p140-positive adult DM associated with hallmark cutaneous disease. Screening of larger IIM cohorts will further characterise the anti-p140 clinical phenotype in comparison to other IIM serological subsets, specifically novel specificities anti-SAE and anti-p155/140.

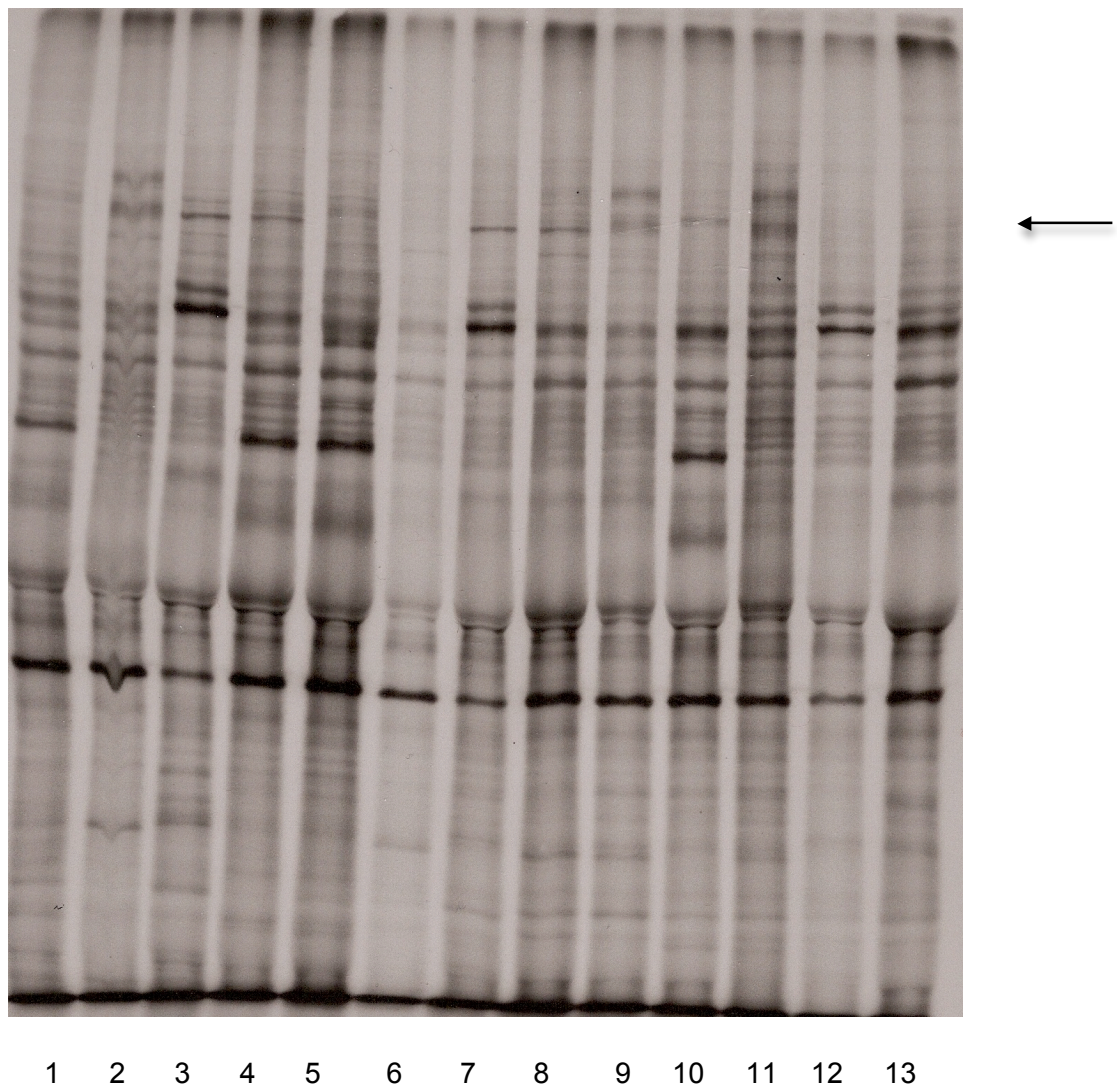
**Figure 22: Immunoprecipitation of anti-p140 autoantibodies in adult DM**

Autoradiogram of 10% SDS-PAGE of immunoprecipitates of [<sup>35</sup>S] labelled K562 cell extract using normal serum (lane 1), reference anti-p155/140-adult DM serum (lane 2), R18883 anti-p140-positive serum (lane 3), AOMIC M203 anti-p140-positive serum (lane 4), AOMIC M238 serum – unknown p140-142 doublet (lane 5), AOMIC M260 serum – faint band at 142 kDa (lane 6), AOMIC M352 serum – negative (lane 7), AOMIC M86 serum – weak unknown p140-142 doublet (lane 8), AOMIC M89 anti-p140-positive serum (lane 9), AOMIC M136 anti-p140-positive serum (lane 10), AOMIC M145 anti-p140-positive serum (lane 11), AOMIC M212 anti-p140-positive serum (lane 12), AOMIC M347 serum – negative (lane 13), and R23389 anti-p140-positive serum (lane 14). Arrow indicates 140 kDa band.



**Figure 23: Immunodepletion experiments with reference anti-p140-positive JDM and adult DM sera, and anti-p155/140-positive JDM sera**

Autoradiogram of 10% SDS-PAGE of immunoprecipitates using anti-p140-positive JDM serum with p140 predepleted [ $^{35}\text{S}$ ] labelled cell extract (using reference R18883 anti-p140-positive adult serum) (see corresponding negative lane 13) / using R23389 anti-p140-positive adult DM serum with p140 predepleted  $^{35}\text{S}$ -labelled cell extract (using reference R18883 anti-p140-positive adult serum) (see corresponding negative lane 12) / using reference anti-p155/140-positive adult DM serum with p140 predepleted [ $^{35}\text{S}$ ] labelled cell extract (using reference R18883 anti-p140-positive adult serum) (see corresponding p155/140 bands lane 11). Arrow indicates 140 kDa band.



**Table 23: Anti-p140 autoantibody ANA IIF patterns (adult DM)**

Sample	ANA IIF pattern
R18883	Fine speckle nucleolar sparing and fine cytoplasmic speckle
R23389	Fine speckle nucleolar sparing 1/40
M89	Homogeneous
M136	Fine speckle nucleolar sparing
M145	Homogeneous
M203	Fine speckle nucleolar sparing 1/40
M212	Homogeneous

**Table 24: Clinical Features of anti-p140 autoantibodies in adult dermatomyositis**

Clinical Feature	Anti-p140 autoantibodies	
	Positive (n=13)	Negative (n=430)
Gender	53.8% Female	74.2% Female
Gotttron's Lesions	92.3%	74.3%
Heliotrope Rash	76.9%	66.8%
Raised Creatinine Kinase	53.9%	76.4%
Systemic Features	76.9%	53.4%
Cancer	0%	14.9%
Interstitial pneumonia	61.5% *	26.2%
Weakness	69.2%	88.3%

\* $P_{uncorr}$ =0.016 (OR 4.5 95% CI: 1.4-14.4)

## **7.2 Summary and Final Discussion**

### **7.2.1 Myositis-specific autoantibodies**

The work described in this thesis combined with various studies over the past few years provides evidence highlighting the importance of autoimmunity in myositis. Up to 80% adult patients and 60% of juvenile patients now have well defined MSAs and MAAs when tested by detailed serological techniques.

- Eight anti-synthetase autoantibodies collectively define the anti-synthetase syndrome, characterised by myositis, skin disease including mechanic's hands and Gottron's papules, non-erosive arthritis, fever, Raynaud's and interstitial pneumonia. Lung involvement is a major complication and may be the predominant clinical manifestation.
- Anti-SRP autoantibodies define patients severe acute / subacute necrotising myopathy, which may be refractory to standard treatment.
- Anti-Mi-2 autoantibodies define patients with classic hallmark DM who appear to respond well to standard treatments and run a monocyclic disease course.
- Anti-CADM-140 (MDA5) autoantibodies appear to be unique to Asian clinically-amyopathic DM patients with a high risk of progressive acute interstitial pneumonia.
- Anti-p155/140 is a novel autoantibody found in both adult DM and JDM. This specificity defines patients with severe cutaneous disease, and is associated with cancer-associated myositis in older adults.
- Anti-SAE autoantibodies are detected in adult DM who may present with clinically-amyopathic DM first, progressing to myositis and systemic features. Interstitial lung involvement appears infrequent.
- Anti-p140 autoantibodies are a major serological subset in JDM, and are associated with calcinosis.

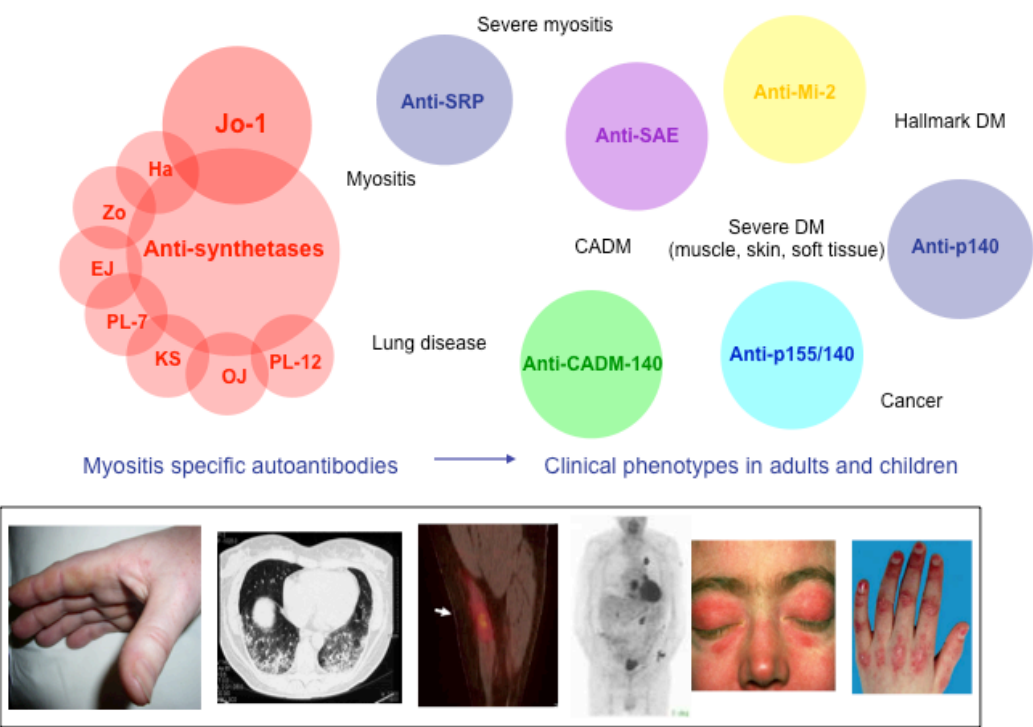
The identification and characterisation of an increasing number of MSAs and their corresponding targets has emphasised how we should now consider a clinico-serological classification in IIM, rather than simply diagnose patients with PM, DM or CTD-myositis overlap. In the future, if MSAs are used to define patients into clinical syndromes this may help predict outcomes and thus influence treatment strategies.

**Table 25: Summary: Myositis-specific autoantibodies, autoantigen targets and clinical features**

Autoantibodies	Target autoantigen and function	Clinical Phenotype	Autoantibody frequency (%)	
			Adults	JDM
Anti-ARS	ARS – intracytoplasmic protein synthesis	Anti-synthetase syndrome	30-40%	1-3%
Anti-Jo-1	Histidyl	Myositis, mechanic's hands, Gottron's papules, arthritis, fever, Raynaud's phenomenon, high frequency of interstitial pneumonia		
Anti-PL-7	Threonyl			
Anti-PL-12	Alanyl			
Anti-EJ	Glycyl			
Anti-OJ	Isoleucyl			
Anti-KS	Asparaginy			
Anti-Ha	Tyrosyl			
Anti-Zo	Phenylalanyl			
Anti-SRP	SRP – intracytoplasmic protein translocation (6 polypeptides and ribonucleoprotein 7SLRNA)	Acute onset necrotizing myopathy (severe weakness, high CK) May be refractory to treatment	5%	<1%
Anti-Mi-2	Helicase protein – nuclear transcription (Forms the NuRD complex)	Adult DM and JDM (hallmark cutaneous disease, milder muscle disease with good response to treatment)	<10%	<10%
Anti-p155/140	TIF1-γ (p155) – nuclear transcription + cellular differentiation	CAM in adult DM Severe cutaneous disease in adult DM and JDM	13-21%	23-29%
Anti-p140	Likely to be NXP-2 – nuclear transcription + RNA metabolism	JDM with calcinosis	N/A	23%
Anti-SAE	SAE - post-translational modification (targets include transcription factors)	Adult DM May present with CADM first	5%	N/A
Anti-CADM-140	Intracytoplasmic MDA5 – innate immune responses against viral infections	CADM Rapidly progressive interstitial pneumonia	Overall - unknown	N/A



**Figure 24: Summary: Myositis-specific autoantibodies and clinical features**



### 7.2.2 Pathogenic mechanisms: MSAs and autoantigens as clues

There is strong evidence that autoimmune mechanisms play an important role in both polymyositis (PM) and dermatomyositis (DM). Previous studies have reported that around 40-50% of adults and 20-30% of juvenile IIM patients have the presence of high-titre autoantibodies. Myositis autoantibodies can be categorised into myositis associated autoantibodies (MAAs) and myositis specific autoantibodies (MSAs). The MAAs; anti-U1-RNP, anti-U3-RNP (fibrillarin), anti-PM-Scl and anti-Ku are principally seen in myositis-scleroderma overlap syndromes (240), whereas the MSAs are highly selective, mutually exclusive and are associated with particular genotypes and clinical phenotypes within the myositis spectrum (6, 8, 9, 241). This thesis describes a comprehensive, clinical and serological study of adult and juvenile idiopathic inflammatory myopathy. Based on this body of work combined with additional preliminary studies, the frequency of MSAs/MAAs is nearer 80% in adults and 60% in children, therefore dispelling the notion that the majority of myositis cases are 'seronegative' (241-243).

MSAs are directed against both cytoplasmic and nuclear components of the cell involved in key regulatory intra-cellular processes, including gene transcription, protein translocation and anti-viral responses. Interestingly, autoantigens with analogous cellular functions are associated with similar clinical subsets. For example, as described in previous chapters, the anti-ARS autoantibodies target functionally related cytoplasmic tRNA enzymes involved in protein synthesis. This group of autoantigens form the largest subset recognised in adult IIM patients, and are associated with distinct clinical features collectively termed the anti-synthetase syndrome. A further subset of myositis patients with severe necrotising myopathy is characterized by the presence of autoantibodies directed against the signal recognition particle (SRP) (121, 122). SRP is a cytoplasmic ribonucleoprotein complex that recognizes secreted and membrane-bound proteins regulating protein translocation through the endoplasmic reticulum.

In contrast, autoantibodies associated with DM clinical phenotypes, such as anti-Mi-2 and novel MSAs anti-p155/140 and anti-p140, target nuclear proteins (126, 143, 157, 158, 161). Mi-2 is a nuclear helicase protein that forms part of the nucleosome remodelling deacetylase (NuRD) complex playing a role in gene transcription, specifically chromatin remodelling (129). Targoff *et al* has identified the p155 target of the p155/140 complex as transcriptional intermediary factor 1-gamma (TIF1-γ), a nuclear protein involved in cellular differentiation (159). In addition, the p140 target



has been identified as nuclear matrix protein NXP-2, which plays a role in RNA metabolism and maintenance of nuclear architecture (228, 234). Finally, anti-SAE autoantibodies target the small ubiquitin-like modifier activating enzyme subunits involved in post-translational modification that is located in both the nucleus, but unlike other DM-associated autoantigens, it is also found in the cytoplasm of cells (168). The potential relationship between the DM-specific autoantigens is thus of interest. It is possible SAE plays a central role forming stable conjugates with other proteins including transcription factors (168). Interestingly, a specific target of sumoylation is a protein (p66) that has been shown to mediate transcriptional repression of the Mi-2/NuRD complex (178). Furthermore, NXP-2 has been reported to have a role in sumoylation-mediated regulation of transcription (235). Post-translational modification can lead to the generation of self-antigens and so this observation suggests shared pathogenic mechanisms.

The exception to the above observations may be the anti-CADM-140 autoantibody, now described in two Asian DM cohorts with CADM and severe lung involvement (130, 131, 244). The identity of this cytoplasmic protein CADM-140 has recently been discovered to be melanoma-differentiation-associated gene 5 (MDA5) (131, 244). The MDA5 autoantigen appears to be distinct in terms of its biological function and disease expression. MDA5 is involved in the innate immune defence against viral infections through the detection of viral dsDNA (132). This observation may be explained by the fact that anti-CADM-140-autoantibody DM appears to be a unique clinical subset found only in Asian ethnicity, with specific genetic and environmental determinants.

In general, the systemic autoimmune diseases are a genetically complex heterogeneous group of diseases in which the immune system attacks diverse but highly specific intracellular self-structures. In IIM, the striking association between specific autoantigen targets and clinical phenotypes, to the degree that IIM can now be classified into more homogeneous subsets, suggests that the autoantigen itself may be central in determining disease expression. Moreover, in IIM there are key target tissues, in particular the muscle, the skin, the lungs, and in some cases tumour tissue. It is possible that in IIM syndromes, disease initiation is dependent on the structure and concentration of autoantigens (i.e. novel non-tolerized forms – truly novel epitopes or previously recognised but modified epitopes that lower the threshold for autoreactive T cells to be activated) in certain localised pro-immune microenvironments. Several studies have provided insight not only into the potential

importance of the autoantigen target but also its expression in specific microenvironments. Two studies have highlighted how certain autoantigenic -tRNA synthetases (histidyl, asparaginyl and tyrosyl) have chemoattractant properties and can induce leukocyte migration via the CCR5 and CCR3 receptors respectively (135, 245). In comparison, non-antigenic -tRNA synthetases (aspartyl and lysyl) do not activate chemokine receptors. In addition, mononuclear cells expressing chemokine receptors are present in myositis muscle but not normal muscle (135). The authors suggested that autoantigenic ARS are over-expressed in damaged muscle cells and their pro-inflammatory properties promote the immune response, which leads to the development of myositis.

Several studies have now demonstrated that certain autoantigens are enriched in lesional tissue involved in IIM. In the seminal paper by Casciola-Rosen *et al* several myositis-specific and associated autoantigens (including the Jo-1 and Mi-2 proteins) were shown to be upregulated in myositis muscle in comparison to normal muscle especially in regenerating muscle cells, rather than mature myotubes (107). Zampieri *et al* demonstrated similar findings in newborn skeletal muscle, in the absence of infiltrating inflammatory cells. Using immunocolocalisation techniques, anti-Jo-1 sera showed a positive reaction within the cytoplasm of tibialis anterior muscles (TA) from newborn rat cryosections and anti-Mi-2 positive serum gave a homogenous staining of myonuclei. In contrast, sections of TA from adult rats gave negative staining with both anti-Jo-1 and anti-Mi-2 sera. In addition, staining of Jo-1 and Mi-2 paralleled the staining with anti-MHCemb, further indicating that expression of Jo-1 and Mi-2 is enhanced in myotubes (246). Collectively, these findings support the hypothesis that the presence of candidate myositis autoantigens during reparative myogenesis can drive induction and propagation of the autoimmune response.

It has been proposed that susceptibility to the serine protease, granzyme B, is strongly predictive of autoantigen status. Casciola-Rosen *et al* have shown that isoleucyl-, histidyl- and alanyl-tRNA synthetases, Mi-2 and SRP72 are all cleaved by granzyme B, leading to the release of unique fragments that contain autoantigenic epitopes (104). Work by Levine *et al*, has emphasized the potential pathogenic role of proteolytic cleavage of autoantigens in lesional tissues. They demonstrated that the histidyl-tRNA protein exists in two conformations and whilst there are similar overall expression levels of Jo-1 in different tissues, the granzyme B cleavable form is enriched in the alveolar epithelial layer of lung (108). Therefore, the theory that

distinct microenvironments may shape disease expression was emphasised with this finding that a novel conformation of Jo-1 is enriched in the lung in comparison to other tissues including muscle. This has led to the suggestion that the initiating target tissue for the autoimmune response in the anti-Jo-1 syndrome is the lung with secondary attack to muscle.

Further evidence implicating autoantigen driven responses was suggested in a study by Katsumata *et al* (133). They generated an antigen-induced model of IIM following immunisation of congenic mice with either human or murine Jo-1 protein or amino terminal peptides of Jo-1 protein. Early antibody responses demonstrated strong species-specificity with limited cross recognition of Jo-1 between species. Studies on serum samples 8 weeks after immunisation revealed epitope spreading with the breakdown of B-cell tolerance to ubiquitously expressed native target autoantigens. The presence of anti-Jo-1 IgG isotypes indicated class switching and T-cell dependant B-cell responses, likely reflecting a breakdown in T-cell tolerance to self Jo-1. Assessments of mice after murine Jo-1 immunization demonstrated the presence of skeletal muscle inflammation and interstitial lung disease, which are clinical characteristics of the ASS clinical phenotype in humans.

Work on the Mi-2 protein gives further emphasis to the potential role of target proteins in disease expression. Casciola-Rosen *et al* have previously demonstrated that Mi-2 is preferentially expressed in DM muscle rather than PM muscle, supporting the association between DM and Mi-2 (107). Using a conditional gene targeting approach in a mouse model, Kashiwagi *et al* demonstrated that Mi-2 is essential for the development and repair of the basal epidermis (134). Furthermore, studies by Burd *et al* have reported that following UV radiation, Mi-2 protein levels in a human keratinocyte cell line are rapidly up-regulated and maintained by an increase in translation efficiency through a regulatory element in the 5'-UTR region of mRNA and increased protein stability. This suggests enhanced expression of Mi-2 in UV exposed keratinocytes leads to the initiation of autoimmunity and subsequently the DM phenotype associated with classic skin lesions. Autoantibodies to other members of the NURD complex have not been reported in IIM, and the effects of UV radiation do not significantly affect other NURD proteins, leading to the hypothesis that Mi-2 may act independently of the NURD complex in response to DNA damage (247).

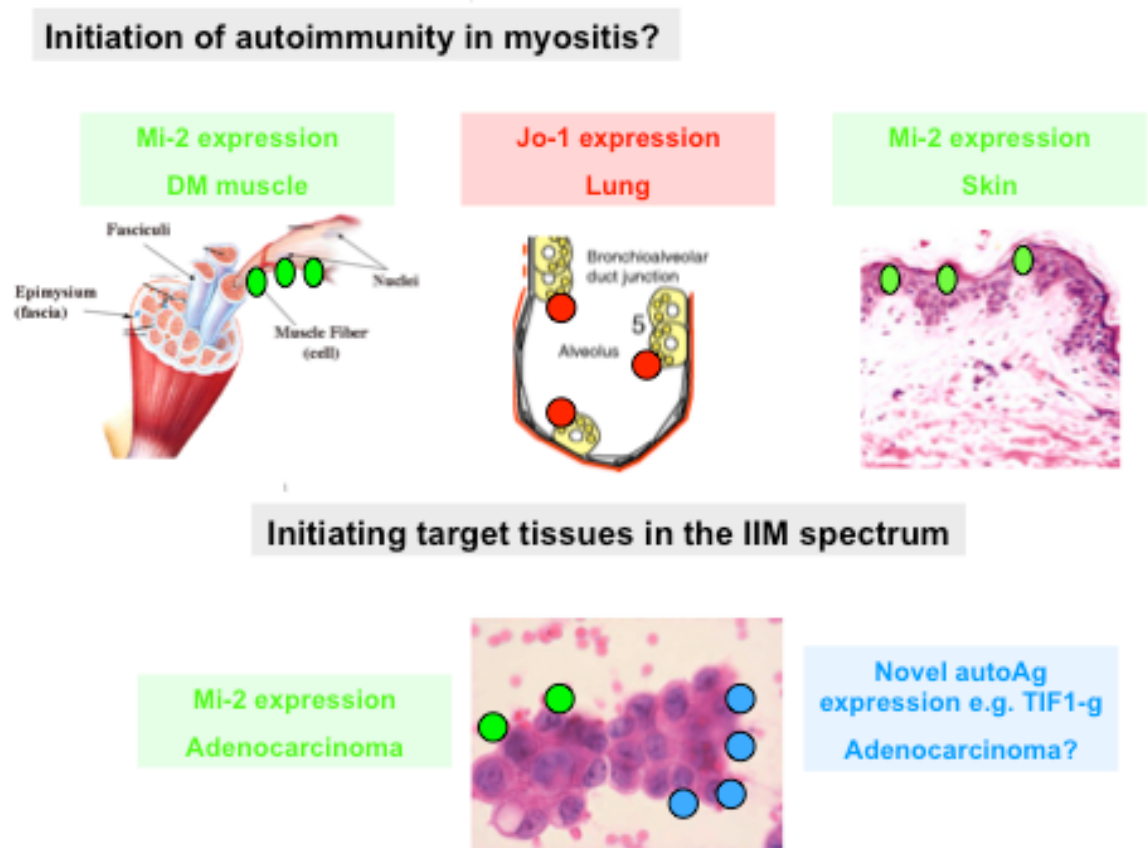
The association between malignancy and myositis, particularly in adult DM is well recognised. Casciola-Rosen *et al* have demonstrated that Mi-2 and Jo-1 expression is increased in certain tumours including breast and lung adenocarcinomas (107). Okada *et al* also demonstrated the presence of autoantibodies to the DNA mismatch repair protein PMS1 (248), a further myositis autoantigen in patients with pancreatic adenocarcinoma, with protein levels increased in the corresponding lesional tissue (249). Furthermore, it has been demonstrated that tumour cells express low levels of specific effector caspases (250, 251), leading to the suggestion that with certain forms of cancer there may be a relative increase in other proteolytic enzymes, including granzyme B. It is therefore plausible that there may be generation of uniquely cleaved autoimmune fragments in carcinoma cells. Suber *et al* have suggested a hypothesis that in certain circumstances, the autoimmune response in myositis is secondary to a primary anti-cancer response. A further trigger, such as a muscle-trophic virus then leads to enhanced expression of autoantigens promoting activation of autoreactive lymphocytes and tissue damage (138). This theoretical model is supported by Mimuro *et al*, who demonstrated that the cancer-associated anti-p53 antibody is also observed in PM and DM patients, even when not associated with malignancy (252). The discovery of the p155/140 autoantigen and its clear association with cancer in adult DM is an exciting breakthrough, and may increase our understanding of the relationship between autoimmunity and cancer (143, 157, 158). What is fascinating is the fact that the same protein is an autoimmune target in JDM, which does not appear to be associated with malignancy (see Chapter 6) (161), and this also appears to be the case in younger adults with amyopathic DM (as described in Chapter 4). Perhaps some perturbation of p155/140 in proliferating cells combined with a more efficient anti-cancer response by a younger immune system may be the answer. Interestingly p155 (TIF1- $\gamma$ ) has been shown to inactivate Smad-4, which regulates TGF- $\beta$  signalling, thus promoting cell growth and differentiation (including malignant tumours) (253). Further studies to investigate p155/140 autoantigen expression and conformation in cancer tissue and compare this to levels in other lesional tissues may be revealing. Whilst the anti-p155/140 autoantibody is the only MSA significantly associated with cancer there is evidence for roles of other autoantigen targets in malignant processes. These include autoantigenic members of the aminoacyl tRNA synthetase family; tyrosyl-, isoleucyl-, phenylalanyl- and glycyl-tRNA synthetase. Of particular interest is the preferential expression of the interstitial lung disease associated autoantigen, phenylalanyl-tRNA synthetase (alpha-subunit) in solid lung tumours and acute myeloid leukaemia (254-260). In addition, MORC3 (NXP-2 / p140) is essential for

regulating the sub-nuclear localisation of p53 (261), and MDA5 activates I $\kappa$ B (IKK) related kinases, known to have a role linking chronic inflammation and cancer (262). The next question of whether MSAs are simply epiphenomena or directly linked to pathogenesis remains uncertain. Two recent publications have highlighted the potential role of anti-Jo-1 autoantibodies in disease activity and pathogenesis. Stone *et al* investigated the association between anti-Jo-1 levels and myositis disease activity (263). The authors performed a cross-sectional study of 81 anti-Jo-1 positive patients and disease activity was assessed using CK levels and the myositis disease activity assessment tool. Anti-Jo-1 levels were measured using a commercial ELISA and a custom ELISA using recombinant human autoantigen. Anti-Jo-1 levels showed modest correlation with CK, myositis and joint disease activity. Longitudinal data was available on 11 patients that showed serial anti-Jo-1 levels did correlate with CK and disease visual analogue scales. In the second study, Eloranta *et al* concluded that immune complexes containing either anti-Jo-1 or anti-Ro in the presence of RNA may act as endogenous inducers of type 1 interferon alpha (IFN- $\alpha$ ) (80). Together, these findings suggest that anti-Jo-1 autoantibodies may play a role in disease propagation, and it would be of interest to see whether other MSAs correlate with disease activity and are able to induce pro-inflammatory cytokine activity. Therefore in IIM, disease propagation and/or disease flares may be autoantibody driven in an already primed pro-immune environment, which leads to autoantibody mediated autoantigen capture, presentation and opsonisation.

It now appears clear that in IIM specifically targeted autoantigens share unique properties, are preferentially expressed in disease-associated tissues and have a role in disease initiation and propagation. Investigating the structure and function of target molecules, and whether autoantibodies themselves have functional roles appears critical for understanding the pathogenic mechanisms in this complex spectrum of diseases. This in turn, may lead to therapeutic advances including the development of more targeted treatments.

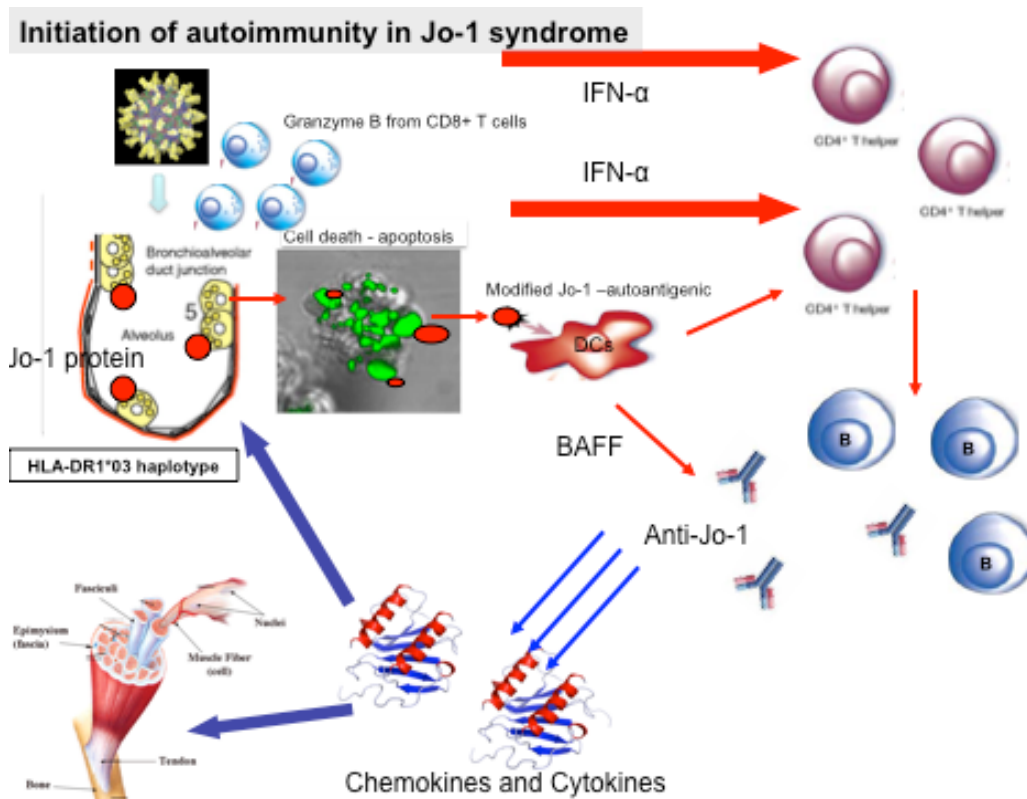
### 7.2.3 Hypothetical models of autoreactivity and pathogenesis in myositis subsets

Figure 25: Summary of observations: autoantigens as clues in target tissues



- The Mi-2 protein is enriched in DM muscle in comparison to PM muscle and normal muscle (107).
- In mouse models, the Mi-2 protein plays a critical role in the development of the skin basal epidermal layer (134).
- In vitro, UV light upregulates Mi-2 expression in human keratinocyte cell lines (264).
- The Mi-2 protein is enriched in adenocarcinoma tissue (107).
- Expression of the Jo-1 protein is highest in the lung in comparison to other tissues including muscle (108).
- The Jo-1 protein is highly susceptible to granzyme B cleavage in the lung and the immunogenic form of Jo-1 in vivo exists unbound (108).
- Mice immunised with the Jo-1 protein develop the muscle inflammation and lung fibrosis synonymous with the anti-synthetase syndrome (133).

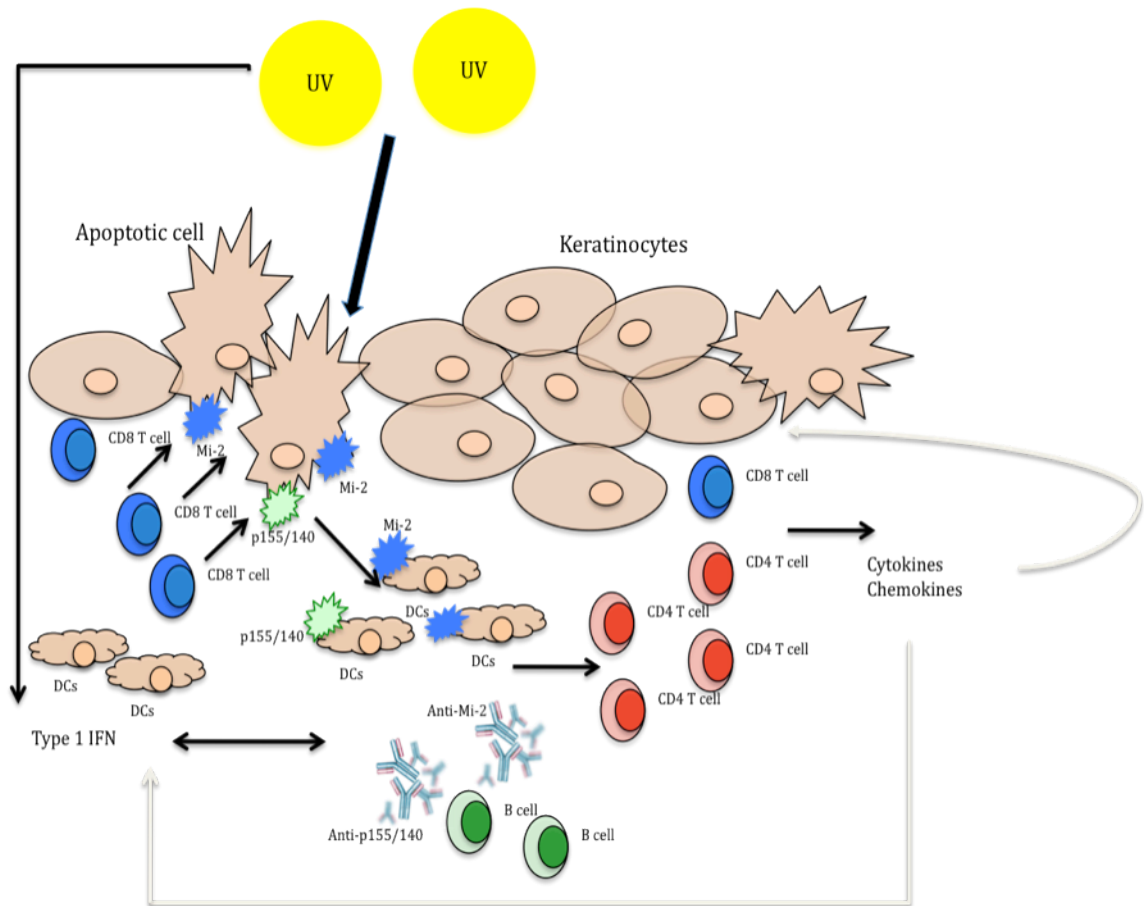
**Figure 26: Lung model**



This model is based on the observation and works of others (80, 103, 104, 106, 108, 135, 137, 265-267). An environmental trigger such as a lung trophic virus and/or an environmental toxin (e.g. smoking leading to damage of the lung epithelial barrier) may initiate the autoimmune response in genetically susceptible individuals (e.g. HLA-DR3 haplotype). Virus particles incorporate into the alveolar epithelial cells triggering cytotoxic CD8 T cell mediated release of the serine protease granzyme B. This leads to alveolar cell apoptosis and thus modification of the Jo-1 protein via granzyme B cleavage. Granzyme B generated fragments of Jo-1 become immunogenic. Jo-1 is chemo-attractant via CCR5 receptor binding, thus attracting CD4/8 T cells, monocytes and immature dendritic cells (DCs). The Jo-1 autoantigen may be incorporated into DCs and then presented to CD4 T cells, which in turn initiates T cell autoreactivity and autoantibody production by B cells. This mechanism is enhanced by anti-viral type 1 IFN responses leading to DCs maturation and further antigen presentation. Anti-Jo-1 autoantibody production is also promoted through the interaction of mature DCs with autoreactive B cells via BAFF. This scenario is supported by the finding that bronchial-alveolar fluid from NSIP lung (the most common subtype of interstitial pneumonia in Jo-1 myositis) is rich in CD8 T cells. Therefore, the initiating target tissue in Jo-1 ASS may be the lung with muscle and/or skin targeted at later stages of injury as Jo-1 autoantigen

levels and corresponding autoantibody production increases. It is likely that the other ARS proteins, in particular PL-7 and PL-12 share similar immunogenic properties and expression profiles in the lung, particularly because lung damage is the predominant feature of this subtype of anti-synthetase syndrome.

**Figure 27: Skin model: Initiation of autoimmunity in Mi-2 dermatomyositis syndrome**



In genetically susceptible individuals (e.g. DRB1\*07 in Mi-2 syndrome), UV-light triggers cutaneous inflammation. CD8 T cell mediated keratinocyte cell death leads to upregulation of unique immunogenic fragments of the Mi-2 protein. It is possible that the Mi-2 autoantigen also has chemoattractant properties (similar to the Jo-1 autoantigen), thus recruiting T helper cells and DCs. Antigen presentation promotes T cell autoreactivity and anti-Mi-2 autoantibody production by B cells. This process is further enhanced by UV-light induced IFN production leading to DCs maturation and endothelial cell activation. Finally, loss of normal Mi-2 function disrupts basal epidermal cell layer repair mechanisms propagating epidermal damage. Similar to the Jo-1 lung model, attack on muscle may be a secondary downstream effect in



this model. The p155/140 protein or other DM-specific autoantigens such as SAE may be modified by similar mechanisms.

**Cancer model: as proposed by Suber *et al* (138)**

It is proposed that perhaps in some patients with cancer-associated dermatomyositis the first event in the pathogenic cascade is the initiation of an anti-tumour response. In the early stages of abnormal cell differentiation, the immune system recognises highly expressed tumour antigens (based on current observations these antigens may include Mi-2 and p155/140). This response is either highly effective leading to efficient removal of tumour cells (before any pathological and clinical manifestations) (perhaps this mechanism may explain why p155/140 is a major autoimmune target in JDM) or ineffective (i.e. development of cancer in an 'older' less efficient immune system). This powerful anti-tumour response combined with other stressors e.g. toxins or trophic viruses leads to muscle or cutaneous injury. For example, muscle cells may undergo apoptosis, and those antigens initially targeted by the anti-cancer response, become upregulated in areas of repair and regeneration. This in turn leads to antigen presentation, autoreactive T cell propagation and autoantibody production. Inflammatory cells and cytokine production promote further injury and thus repair, which sustains the autoimmune response against cancer, muscle and/or skin cells. With the discovery of the p155/140 autoantigen, this hypothesis can now be interrogated.

### **7.3 Future Work**

Future work should focus on increasing our understanding the significance of novel protein targets, from how they are expressed at a cellular level to how they classify patients based upon clinical features.

#### **7.3.1 Screening larger cohorts of IIM patients to further define clinico-serological subsets**

Further screening is required to establish the prevalence and disease specificity of novel autoantibodies in a European-wide cohort of adult patients with IIM. With the formation of the European Myositis Consortium, cohorts of adult patients with IIM with linked clinical data and serum samples are now available for analysis. This combined with patients recruited to the UK AOMIC Registry and UK JDM cohort study will form the largest collection of adult and juvenile IIM patients (over 1000 patients) with comprehensive clinical data (see chapter 2, appendix, proforma 3 and 4). MSAs are associated with clinical phenotypes and the study of a large cohort of patients will form a major biomarkers resource. Rather than diagnose patients with PM or DM, the work described in this thesis and future studies will lead to defining IIM patients into clinico-serological syndromes, which in the future will aid physicians to predict clinical outcomes and plan specific treatment regimes.

#### **7.3.2 Investigate the persistence and level of anti- p155/140 in JDM in relation to increasing age and clinical course**

Around 20% of children with JDM are positive for anti-p155/140 autoantibodies. The same autoantibody specificity is found in adult DM associated with malignancy. Therefore, it will be important to investigate how these autoantibodies fluctuate over disease course (disease activity and response to treatment). Serial samples on children recruited to the UK JDM Cohort Study have been stored during disease course annually will be screened for anti-p155/140 using IPP. Autoantibody levels can be quantified using immunoprecipitation blotting. Briefly, immunoprecipitations will be completed using 40µl serum (p155/140 and control serum), 2mg protein A Sepharose and 5 mM bis-(sulphosuccinimidyl)-suberate cross-linker. Beads will be incubated in 2ml unlabelled K562 cell extract for a total of 2 hr at 4°C. Samples will be resuspended in 80µl SDS sample buffer and are heated. Proteins are fractionated by 10% SDS-PAGE and transferred to nitrocellulose by Western blotting. Blots will be probed with anti-p155/140 positive sera and bands will be detected using an alkaline phosphatase conjugated donkey anti-human IgG antibody and the BCIP/NBT liquid substrate solution. The immunoblots will then be

quantified by densitometry. Equal loading will be confirmed by spiking samples with a pre-defined amount of a lower molecular weight protein. Blots will be stripped and re-probed for this protein. Autoantibody data will be linked to prospective clinical data collected every six months and recorded on the JDRR database.

### **7.3.3 Antigen expression**

#### ***Lesional tissue***

As described in section 7.2.2, based on the paradigm suggested by Casciola-Rosen and Rosen that in IIM lesional tissue is enriched for certain autoantigens that, under certain circumstances, may drive a dysregulated autoimmune response, further work is required to investigate this hypothesis. IIM is an excellent model to explore this concept further. Muscle, lung, skin and cancer biopsy material is readily accessible tissue that can be studied directly. Tissue expression can be investigated by immunohistochemistry - biopsy specimens (stored at -80°C) will be cryostat (6-8 um thick), mounted and dried. Slides will be warmed at room temperature, fixed in ice-cold acetone and air-dried. Sections will be washed in PBS and rinsed in TBS-tween. Sections will be blocked and incubated overnight at 4°C in a humidified atmosphere with commercial monospecific primary antibodies to myositis specific antigens (SRP54 chicken polyclonal ab14072, Mi-2 rabbit polyclonal 06-878, TIF1-γ mouse monoclonal ab33475, SAE1 sheep polyclonal alx210328, SAE2 rabbit polyclonal ap10659, HARS mouse monoclonal ab50835, TARS mouse monoclonal ab50147, ALARS rabbit polyclonal ab50147, NARS rabbit polyclonal ab50144, GARS rabbit polyclonal ab42905, Isoleucyl tRNA-synthetase rabbit polyclonal ab31533, Phenylalanyl tRNA-synthetase alpha H00002193-A01 mouse polyclonal, Phenylalanyl tRNA-synthetase beta H00010056-A01 mouse polyclonal. Sections will be washed and incubated with the appropriate secondary antibodies (anti-mouse/rabbit/sheep/chicken IgG) conjugated with horseradish peroxidase, stained with 1-Step 4-CN (Perbio) and visualised. Negative controls will be performed by omitting or isotype-matching with the primary antibody. Sections will also be probed with FITC and TRITC conjugates as secondary antibodies in order to co-localise autoantigens by double immunofluorescence with appropriate cell types present in lesional tissue. IPP and immunoblotting on cell extracts will be performed to ascertain the specificity of commercially prepared antibodies prior to immunohistochemistry (as already done with commercial anti-SAE1/2 and anti-NXP-2 demonstrating that these antibodies do recognise corresponding antigen targets derived from human cell lines.

### ***Different cell lines***

Although the autoantigen targets can be visualised using  $^{35}\text{S}$ , for characterisation with mass spectrometry it may be necessary to scale up the quantity of polypeptide immunoprecipitated (e.g. if it is constitutively expressed in low copy number). The initial approach will be to increase the quantity of protein-A-Sepharose beads and source of antigen. However, subsequently, a combination of sub-cellular fractionation and immunoaffinity chromatography may be used to optimise antigen recovery. In addition, other human cell lines may contain different levels of expressed known and novel myositis autoantigens. Differences of expression in these seemingly ubiquitously expressed proteins will provide additional insight into pathogenic mechanisms.

#### **7.3.4 Development of quantitative solid-phase assays**

The future development of commercial assays that test for MSAs in routine clinical practice is important. To date, certain myositis autoantigens (Jo-1, PL-7, PL-12, Mi-2 and SRP) have been expressed as recombinant proteins for use in commercial ELISAs and line-blot. An important aim is to develop further assays for the novel MSAs including the detection of anti-p155/140 (p155 autoantigen - TIF1- $\gamma$  (159)), given the potential importance of this autoantibody in the diagnosis of JDM and its association with malignancy in adult-onset DM. Furthermore, the development of custom-based ELISAs will enable the quantitative analysis of MSA levels in serum samples that can be correlated with measures of disease activity, as previously demonstrated with anti-Jo-1 autoantibodies (263). Other autoantigen systems that may also become strategically relevant include anti-SAE, anti-p140 and anti-CADM-140. To develop sensitive immunoblotting techniques and custom based ELISAs preparations will be enriched with the appropriate autoantigens by a combination of cell fractionation, HPLC and affinity purification using monoclonal antibodies. An alternative approach is to extract mRNA from cell lines e.g. K562, A549 cells to develop a cDNA library. This will be screened by either PCR or using a SEREX. Positive sequences will be expressed in a suitable system for the large-scale production of the recombinant peptide. In addition, commercial supplies of the relevant autoantigen can be sourced e.g. TIF1- $\gamma$  peptide (AbCam 47108) that will be tested for autoantibody recognition by immunoblotting using a panel of appropriate positive and negative controls. A further approach will be the generation of 20-mer overlapping synthetic peptides spanning the known sequence of autoantigens.

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